

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
1 May 2003 (01.05.2003)

PCT

(10) International Publication Number
WO 03/035846 A2

(51) International Patent Classification⁷: C12N

St., Apt. 408, Denver, Colorado 80206 (US). XU, Liang-
guo; 831 Cherry St., Apt 25, Denver, CO 80220 (US).

(21) International Application Number: PCT/US02/34376

(22) International Filing Date: 24 October 2002 (24.10.2002)

(74) Agents: CONNELL, Gary, J. et al.; Sheridan Ross P.C.,
Suite 1200, 1560 Broadway, Denver, CO 80202-5141 (US).

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:

60/345,106	24 October 2001 (24.10.2001)	US
60/348,962	14 January 2002 (14.01.2002)	US
60/354,966	7 February 2002 (07.02.2002)	US
60/403,364	13 August 2002 (13.08.2002)	US

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VC, VN, YU, ZA, ZM, ZW.

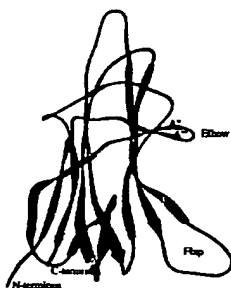
(71) Applicant: NATIONAL JEWISH MEDICAL AND RE-
SEARCH CENTER [US/US]; 1400 Jackson St., Denver,
CO 80206 (US).

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

(72) Inventors: ZHANG, Gongyi; 3635 South Hibiscus Way,
Denver, CO 80206 (US). SHU, Hong-Bing; 4801 9th Ave.,
#209, Denver, CO 80220 (US). LIU, Yingfang; 1150 Vine

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(54) Title: THREE-DIMENSIONAL STRUCTURES OF TALL-1 AND ITS COGNATE RECEPTORS AND MODIFIED PRO-
TEINS AND METHODS RELATED THERETO



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C

(57) Abstract: Disclosed are TALL-1 and TALL-1 receptor protein homologues (agonists and antagonists) designed based on the three-dimensional structure of sTALL-1, eBCMA and eBAFF-R; agonist homologues of APRIL; methods of using wild-type APRIL to inhibit the activity of TALL-1; compositions comprising such homologues, nucleic acid molecules encoding such homologues, and therapeutic methods of using such compounds and compositions. Also disclosed are crystalline complexes of sTALL-1 and sTALL-1 in complex with either BCMA or BAFF-R; models of three-dimensional structures of such crystalline complexes and related structures, methods of drug design using any portion of such structures; methods of design and/or identification of regulatory peptides derived from the such structures; compounds identified by drug design using such structures; and the use of such compounds in therapeutic compositions and methods.



Published:

— without international search report and to be republished
upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

THREE-DIMENSIONAL STRUCTURES OF TALL-1 AND ITS COGNATE RECEPTORS AND MODIFIED PROTEINS AND METHODS RELATED THERETO

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Field of the Invention

This invention generally relates to the three-dimensional structure of sTALL-1 and sTALL-1 in complex with the extracellular domains of its cognate receptors, BCMA and BAFF-R, and to the use of such structures to develop agonists and antagonists and lead compounds for drug development in the area of therapeutic agents related to TALL-1 biological activity.

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Background of the Invention

TNF (tumor necrosis factor) family ligands and their corresponding receptors (TNFR) play pivotal roles in mammalian cell host defense processes, inflammation, apoptosis, autoimmunity, and organogenesis. There are at least 18 TNF ligands and 27 receptors identified so far. Some ligands have multiple receptors, and some receptors also bind multiple ligands. The interactions between ligands and receptors are usually very specific and have high apparent affinity (0.1nM-1nM) (Locksley et al., 2001, *Cell* 104:487-501; Fesik et al., 2000, *Cell* 103:273-282).

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The first TNF ligand trimer structure (TNF α) was determined more than a decade ago (Jones et al., 1989, *Nature* 338:225-228; Eck et al., 1989, *J. Biol. Chem.* 264:17595-17605). It consists entirely of β strands and loops. The structure has a standard 'jellyroll' topology and is remarkably similar to capsid proteins of small RNA viruses such as satellite tobacco necrosis virus (Jones et al., 1984, *J Mol Biol* 177: 735-767). Three monomers of TNF α form a trimer through highly conserved hydrophobic surfaces. The trimer also exists in solution. Structures of TNF β , CD40L, and TRAIL were subsequently determined (Eck et al., 1992, *J. Biol. Chem.* 267:2119-2122; Karpusas et al., 1995, *Structure* 3:1031-1039; Cha et al., 1999, *Immunity* 11:253-261). These structures are similar to the TNF α structure, although the sequence homology is low (20-25%) among the TNF family members. These studies led to proposals that all TNF family members have similar structures and function as trimers (Locksley et al., 2001, *supra*; Fesik, 2000, *supra*). Due to the scarcity of available structures (4 of 18) and low sequence homology among the TNF family members, the generality of this conclusion is unclear.

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The structure of the complex of TNF β and cysteine-rich domains (CRDs) from its cognate receptor, TNFR1, has also been determined (Banner et al., 1993, *Cell* 73:431-445). The structure showed that the three elongated receptor domains bind to one TNF trimer at the interfaces formed between the TNF monomers. Two CRDs (CRD2 and CRD3) make contacts with two distinct regions of TNF β . The recently determined complex structure of TRAIL and DR5 disclosed a similar interaction mode as observed in the TNF β and TNFR1 co-crystal structure, although CRD3 of DR5 assumes a different orientation compared to the one in the TNF β and TNFR1 structure (Mongkolsapaya et al., 1999, *Nat. Struct. Biol.* 6:1048-1053; Hymowitz et al., 1999, *Mol. Cell* 4:563-571). It was proposed that the TNF trimeric ligands trigger the trimerization of their cognate receptors, which causes the cytoplasmic regions of the receptor to form a cluster that can recruit adaptor proteins, leading to the activation of downstream signal transduction pathways (Fesik, 2000, *supra*; Banner, et al., 1993, *supra*; Mongkolsapaya, et al., 1999, *supra*; Hymowitz, et al., 1999, *supra*). This theory is now challenged by new findings showing that TNF receptor and Fas exist in an oligomeric state through the pre-ligand-binding assembly domain (PLAD) before the binding of ligands (Chan et al., 2000, *Science* 288:2351-2354; Siegel et al., 2000, *Nat. Immunol.* 1:469-474).

TALL-1, also known as, BAFF, THANK, BLyS and zTNF4, and its receptors BCMA, BAFF-R and TACI are four recently identified TNF/TNFR (TNF receptor) family members (Shu et al., 1999, *J. Leukocyte Biology* 65:680-683; Schneider et al., 1999, *J Exp Med.* 189:1747-56; Moore et al., 1999, *Science* 285:260-263; Mukhopadhyay et al., 1999, *supra*; Shu et al., 2000, *Pro. Natl. Acad. Sci. USA* 97:9156-9161; Gross et al., 2000, *Nature* 404:995-999; Thompson et al., 2000, *J Exp Med* 192:129-35; Marsters et al., 2000, *Curr Biol.* 10:785-8; Xia et al., 2000, *J Exp Med.* 192:137-43; Yan et al., 2000, *Nat. Immunol.* 1:37-41; Yu et al., 2000, *Nat Immunol.* 1:252-6; Thompson et al., 2001, *Science* 293: 2108-2111; and Yan et al., 2001, *Curr Biol.* 11(19):1547-52). Overexpression of sTALL-1 in mice leads to increased numbers of mature B-lymphocytes, splenomegaly, anti-DNA antibodies, proteinuria, and glomerulonephritis. These phenotypes mimic those of systemic lupus erythematosus (Shu et al., 2000, *supra*; Gross et al., 2000, *supra*; Thompson et al., 2000, *supra*; Marsters et al., 2000, *supra*; Xia et al., 2000, *supra*; Yan et al., 2000, *supra*; Mackay et al., 1999, *J Exp Med* 190:1697-710; Khare et al., 2000, *Proc Natl Acad Sci USA* 97:3370-

5) The experiments of BAFF knock-out showed that BAFF was absolutely required for normal B cell development (Schiemann et al., 2001, *Science* 293:2111-2114; Gross et al., 2001, *Immunity* 15(2):289-302). The phenotype is similar to that caused by BAFF-R deficiency (Thompson et al., 2001, *supra*; Yan et al., 2001, *supra*). In the other hand, the knock-outs of BCMA and TACI did not lead to any severe B cell phenotypes (Xu et al., 2001, *Mol. Cell. Biol* 21:4067-4074; Von Bulow et al., 2001, *Immunity* 14:573-582). Interestingly, APRIL (also called TALL-2), the closest family member of TALL-1, does not bind to BAFF-R (Schiemann et al., 2001, *supra*), although it binds to BCMA and TACI with an affinity similar to sTALL-1 (Yu et al., 2000, *supra*).

10 In contrast to the other receptor family members that have at least three to four CRDs in their extra-cellular domains, BCMA and BAFF-R have only one CRD and TACI has two CRDs (Shu et al., 2000, *supra*; Gross et al., 2000, *supra*; Thompson et al., 2000, *supra*; Marsters et al., 2000, *supra*; Thompson et al., 2001, *supra*). Nevertheless, the overall binding affinities of sTALL-1 with BCMA and TACI (0.1nM-1nM) are similar to those of other family members (Yu et al., 2000, *supra*). Furthermore, as predicted from sequence alignment, the CRDs in BCMA and TACI contain A1 and C2 modules (Gross et al., (2000) *Nature*, 404:995-999), which were two of multiple defined structural motifs that characterize extracellular domains of TNF receptors (Naismith et al., 1998, *TRENDS Biochem. Sci.* 23:74-79). The C2 module was also found in TNF-R1 and Fn14 (Bodmer et al., (2002) *J Biol Chem.*, 275:20632-20637), however the C2 in TNF-R1 is not involved in ligand binding (Naismith et al., 1998, *supra*). The only CRD in BAFF-R that was predicted to be the C2 module initially (Thompson et al., 2001, *supra*; Yan et al., 2001, *supra*) has been termed an unknown module X2 recently (Bodmer et al., 2002, *supra*). It is likely that there are novel interactions among these unique ligand-receptor couples accounting for their high affinity (Liu et al., 2002, *Cell* 108:383-394; Bodmer et al., 2002, *supra*). Therefore, to begin to understand the structure and function relationship of TALL-1 and its receptors, and to take advantage of this information to design valuable therapeutic tools, it is necessary to determine the crystal structure of sTALL-1 and its receptors.

Summary of the Invention

One embodiment of the invention relates to a TALL-1 antagonist protein, wherein the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one
5 modification in the region connecting β strands D and E that reduces the biological activity of the TALL-1 antagonist as compared to wild-type TALL-1. In one aspect, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from Val217, His218, Val219, Phe220, Glu221, Asp222,
10 Glu223, and Leu224. In one aspect, the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least two amino acid residues selected from Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224. In another aspect, the TALL-1 antagonist protein comprises an amino acid sequence that differs
15 from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least between about 3 and 8 amino acid residues selected from Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224. In yet another aspect, TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of
20 SEQ ID NO:2, by at least a deletion of the following amino acid residues: Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224; in one aspect of this embodiment, the TALL-1 antagonist protein further comprises a substitution of at least one non-natural amino acid residue for the deleted residues.

In one aspect, the above-described TALL-1 antagonist protein has a reduced ability
25 to form a trimer with other TALL-1 monomers. In another aspect, the protein, when in a trimer with two other TALL-1 monomers, reduces the ability of the trimer to interact with other TALL-1 trimers. The two other TALL-1 monomers can be selected from: a wild-type TALL-1 monomer and a TALL-1 antagonist protein, as well as mixtures thereof.

In one aspect, the above-identified TALL-1 antagonist protein binds to a TALL-1
30 receptor selected from BCMA, BAFF-R and TACI. In one embodiment, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an

amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor. In one aspect, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification in at least one amino acid residue selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222. In this aspect, the additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor (e.g., BCMA, BAFF-R and TACI).

Another embodiment of the invention relates to a TALL-1 antagonist protein that comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one modification that reduces interaction between a first trimer and a second trimer. In this embodiment the first trimer comprises (a) a monomer of the TALL-1 antagonist protein; and (b) two monomers selected from: wild-type TALL-1 monomers, the TALL-1 antagonist protein monomers, and mixtures thereof. The second trimer comprises monomers selected from wild-type TALL-1 monomers, the TALL-1 antagonist protein monomers, and mixtures thereof. In one embodiment, the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue located in a region of TALL-1 selected from β strand C, β strand F, and the region connecting β strand D to β strand E. In another embodiment, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Ile150, Leu169, Phe172, Tyr192, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Ile250, Lys252, and Glu254. In another embodiment, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Val217, His218, Val219,

Phe220, Glu221, Asp222, Glu223, and Leu224. In another embodiment, protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Tyr192, Lys252, Glu254, His218, Lys216, Glu223, Leu224, Val227, Leu229, Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172. In another embodiment, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Tyr192, Lys252, Glu254, and His218. In yet another embodiment, protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Lys216, Glu223, Leu224, Val227, and Leu229. In another embodiment, protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from: Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172.

Preferably, the TALL-1 antagonist protein binds to a TALL-1 receptor selected from BCMA, BAFF-R and TACI. In one aspect, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor. In one aspect, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification in at least one amino acid residue selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, wherein the additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor (e.g., BCMA, BAFF-R and TACI).

In one embodiment, the above-described TALL-1 antagonist protein has a reduced ability to form a trimer with other TALL-1 monomers.

Yet another embodiment of the present invention relates to a TALL-1 antagonist protein that comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue selected from: Phe194, Tyr196, Tyr246, Leu282, Gln144 and Leu285. In one aspect, the protein has a reduced ability to form a trimer with other TALL-1 monomers. In another aspect, the TALL-1 antagonist protein binds to a TALL-1 receptor selected from BCMA, BAFF-R and TACI. In one embodiment, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor. In another embodiment, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification in at least one amino acid residue selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, wherein the additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor (e.g., BCMA, BAFF-R and TACI).

Yet another embodiment of the present invention relates to a TALL-1 antagonist protein, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue that reduces the biological activity of the antagonist protein as compared to a wild-type TALL-1, wherein the amino acid residue is selected from: Gln144, Ile150, Leu169, Phe172, Tyr192, Phe194, Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285. The amino acid sequence of the TALL-1 antagonist further differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type

TALL-1 and the TALL-1 receptor, wherein the amino acid residue is selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222.

Yet another embodiment of the present invention relates to a TALL-1 antagonist
5 protein, wherein the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, wherein the TALL-1 antagonist protein has reduced binding to a receptor for TALL-
10 1 as compared to wild-type TALL-1. In one aspect, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from: Tyr163, Leu211, Ile233, Pro264, and Leu200. In another aspect, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid
15 sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from: Tyr206 and Leu240. In another aspect, the protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from: Arg265, Glu266 and Glu238. In another aspect, the
20 protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from: Asp222, Asp 273 and Asp275. In another aspect, the TALL-1 receptor is selected from BCMA, BAFF-R, and TACI.

In one embodiment, the above-identified TALL-1 antagonist protein has reduced
25 ability to bind to at least two of BCMA, BAFF-R and TACI. In one aspect, the TALL-1 antagonist protein has reduced ability to bind to each of BCMA, BAFF-R and TACI.

Another embodiment of the invention relates to a composition comprising any of the above-identified TALL-1 antagonist proteins.

Another embodiment of the invention relates to an April agonist protein, wherein the
30 protein comprises an amino acid sequence that differs from SEQ ID NO:4 by at least one modification that increases the binding affinity between the APRIL agonist protein and an

APRIL receptor, as compared to the binding affinity between wild-type APRIL and the APRIL receptor. In one aspect, the protein comprises an amino acid sequence that differs from SEQ ID NO:4 by a modification in at least one amino acid residue selected from: Val133, Thr177, Val181, Ile197, Pro230, Leu58, Tyr96, Phe176, Arg206, and Arg265, wherein the modification increases the binding affinity between the APRIL agonist protein and an APRIL receptor, as compared to the binding affinity between wild-type APRIL and the APRIL receptor. In another aspect, the APRIL receptor is selected from BCMA and TACI. In one embodiment, the at least one modification results in binding of the APRIL to BAFF-R.

Another embodiment of the invention relates to a composition comprising any of the above-identified APRIL agonist proteins.

Another embodiment of the invention relates to a method to inhibit TALL-1 biological activity in a mammal, comprising administering to the mammal any of the above-identified TALL-1 antagonist or APRIL agonist proteins. In one aspect, the protein is a competitive inhibitor of wild-type TALL-1 for binding to a TALL-1 receptor. In another aspect, the mammal has, or is at risk of developing, a disease or condition associated with hyperactive B cell development or B cell hyperproliferation. In one aspect, the mammal has, or is at risk of developing, a disease or condition characterized by increased numbers of mature B-lymphocytes, splenomegaly, anti-DNA antibodies, proteinuria, or glomerulonephritis. In one aspect, the disease is systemic lupus erythematosus.

Yet another embodiment of the invention relates to a recombinant nucleic acid molecule comprising a nucleic acid sequence encoding the amino acid sequence of any one of the above-identified TALL-1 antagonist or APRIL agonist proteins, operatively linked to a transcription control sequence. Yet another embodiment of the invention relates to a method to inhibit TALL-1 biological activity in a mammal, comprising administering to the mammal any of such recombinant nucleic acid molecules, wherein the protein is expressed by a host cell in the mammal. In one embodiment, the protein associates with wild-type TALL-1 monomers expressed by the cell to produce TALL-1 trimers containing the protein with reduced TALL-1 biological activity, as compared to a trimer of wild-type TALL-1 monomers. In another aspect, the protein associates with wild-type TALL-1 monomers

expressed by the cell to produce TALL-1 trimers containing the protein with reduced ability to bind to a TALL-1 receptor, as compared to a trimer of wild-type TALL-1 monomers.

Another embodiment of the present invention relates to a BCMA antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:6 by a modification in at least one amino acid residue selected from: Tyr13, Asp15, Leu17, Leu18, His19, Ile22, Leu26, Arg27, and Pro34, wherein the BCMA antagonist has an increased binding affinity for TALL-1 as compared to wild-type BCMA. In one aspect, the amino acid residue is selected from Leu17 and Leu18. In another aspect, the amino acid residue is selected from Ile22 and Leu26. In another aspect, the amino acid residue is selected from Asp15, Arg27 and Tyr13. In another aspect, the amino acid residue is His19. In another aspect, the amino acid residue is selected from Tyr13, Leu17, Leu18 and Ile22. In another aspect, the amino acid residue is substituted with an amino acid residue selected from: Ile, Met, Phe or Tyr. In one aspect the BCMA antagonist is a soluble protein.

Yet another embodiment of the present invention relates to a BAFF-R antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:8 by a modification in at least one amino acid residue selected from: Asp26, Leu28, Val29, Arg30, Val33, Leu37, Leu38, and Arg42, and Pro45, wherein the BAFF-R antagonist has an increased binding affinity for TALL-1 as compared to wild-type BAFF-R. In one aspect, the amino acid residue is selected from Leu28 and Val29. In another aspect, the amino acid residue is selected from Val33, Leu37, Leu38 and Pro45. In another aspect, the amino acid residue is selected from Asp26 and Arg 42. In another aspect, the amino acid residue is selected from Arg30. the amino acid residue is selected from Leu28, Val29 and Val33. In another aspect, the amino acid residue is substituted with an amino acid residue selected from: Ile, Met, Phe or Tyr. In another aspect, the BAFF-R antagonist is a soluble protein.

Another embodiment of the invention relates to a method to inhibit TALL-1 receptor biological activity in a mammal, comprising administering to the mammal any of the above-identified BCMA or BAFF-R antagonists. In one aspect, the antagonist is a competitive inhibitor of a wild-type TALL-1 receptor for binding to TALL-1.

Yet another embodiment of the invention relates to a method to inhibit the biological activity of TALL-1, comprising administering to a cell that expresses TALL-1 a recombinant

nucleic acid molecule comprising a nucleic acid sequence encoding APRIL, or a biologically active fragment thereof.

Another embodiment of the invention relates to an isolated BAFF-R antagonist, wherein the BAFF-R antagonist consists essentially of the amino acid sequence represented by SEQ ID NO:9, or homologues thereof with substantially the same biological activity.

Yet another embodiment of the invention relates to a method to identify a compound that is a competitive inhibitor of TALL-1 binding to its receptor. The method includes the steps of: (a) contacting a TALL-1 receptor or a TALL-1 binding fragment thereof with a homologue of a TALL-1 protein, wherein the homologue comprises an amino acid sequence with a modification in at least one amino acid residue selected from Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, and Glu238; and (b) detecting whether the homologue binds to the TALL-1 receptor or fragment thereof. Homologues that bind to the TALL-1 receptor or fragment thereof potential competitive inhibitors for binding of wild-type TALL-1 to its receptor. In one aspect the method further includes a step (c) of detecting whether homologues that bind to the TALL-1 receptor or fragment thereof in (b) have a TALL-1 biological activity selected from: an ability to activate signal transduction in the TALL-1 receptor, an ability to form a trimer with two other TALL-1 monomers, an ability to form a trimer with TALL-1 two other TALL-1 monomers that is capable of interacting with other TALL-1 trimers. Homologues that have a decreased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 antagonists, and wherein homologues that have an increased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 agonists. In one aspect, step (b) further comprises comparing the binding affinity the homologue to the TALL-1 receptor or fragment of thereof to the binding affinity of wild-type TALL-1 and the TALL-1 receptor, and the method further comprises step (d) of selecting homologues which have an increased binding affinity to the TALL-1 receptor or fragment of and a decreased TALL-1 biological activity.

Yet another embodiment of the invention relates to a method of structure-based identification of compounds which potentially bind to TALL-1, comprising: (a) obtaining atomic coordinates that define the three dimensional structure of TALL-1; and (b) selecting candidate compounds for binding to the TALL-1 by performing structure based drug design with the structure of (a), wherein the step of selecting is performed in conjunction with

computer modeling. The atomic coordinates are selected from: (i) atomic coordinates determined by X-ray diffraction of a crystalline TALL-1; (ii) atomic coordinates selected from: (1) atomic coordinates represented in any one of Tables 2-12; (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by the atomic coordinates of (1); and (3) atomic coordinates in any one of Tables 2-12 defining a portion of the TALL-1, wherein the portion of the TALL-1 comprises sufficient structural information to perform step (b); and/or (iii) atomic coordinates defining the three dimensional structure of TALL-1 molecules arranged in a crystalline manner in a space group P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217 Å.

In one aspect of the above-identified method, the method further comprises a step (c) of selecting candidate compounds of (b) that inhibit the biological activity of TALL-1. In this aspect, the step (c) of selecting can include: (i) contacting the candidate compound identified in step (b) with TALL-1; and (ii) measuring the biological activity of the TALL-1, as compared to in the absence of the candidate compound.

In another aspect of the above-identified method, the method further comprises a step (c) of selecting candidate compounds of (b) that inhibit the binding of TALL-1 to a TALL-1 receptor. In this aspect, the step (c) of selecting can include (i) contacting the candidate compound identified in step (b) with the TALL-1 or a fragment thereof and a TALL-1 receptor or TALL-1 receptor binding fragment thereof under conditions in which a TALL-1-TALL-1 receptor complex can form in the absence of the candidate compound; and (ii) measuring the binding of the TALL-1 or fragment thereof to bind to the TALL-1 receptor or fragment thereof, wherein a candidate inhibitor compound is selected when there is a decrease in the binding of the TALL-1 or fragment thereof to the TALL-1 receptor or fragment thereof, as compared to in the absence of the candidate inhibitor compound. Preferably, the TALL-1 receptor is selected from BCMA, BAFF-R and TACI.

In yet another embodiment, step (b) of selecting comprises identifying candidate compounds for binding to a receptor binding site of the TALL-1 protein, the receptor binding site comprising an amino acid residue selected from Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222.

In another embodiment, step (b) of selection comprises identifying candidate compounds for binding to the TALL-1 such that trimer-trimer interactions between trimers of TALL-1 monomers is inhibited. In this aspect, the step of selecting can include identifying candidate compounds for binding to TALL-1 at a site including an amino acid residue selected from:
5 Gln144, Ile150, Leu169, Phe172, Tyr192, Phe194, Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285.

Another embodiment of the present invention relates to a therapeutic composition comprising a compound that inhibits the biological activity of TALL-1, the compound being
10 identified by the method described above. Another embodiment of the present invention relates to a method to treat a disease or condition that can be regulated by modifying the biological activity of TALL-1, comprising administering to a mammal with such a disease or condition such a therapeutic composition.

Yet another embodiment of the invention relates to a method to construct a three
15 dimensional model of TALL-1 protein or homologue thereof, comprising: (a) obtaining atomic coordinates that define the three dimensional structure of TALL-1, the atomic coordinates being selected from any of those described above; and (b) performing computer modeling with the atomic coordinates of (a) and to construct a model of a three dimensional structure of a TALL-1 or homologue thereof.

20 Yet another embodiment of the invention relates to a method of structure-based identification of compounds which potentially bind to a TALL-1 receptor selected from BCMA and BAFF-R, comprising (a) obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R; and (b) selecting candidate compounds for binding to the BCMA or BAFF-R by performing structure based drug design with the
25 structure of (a), wherein the step of selecting is performed in conjunction with computer modeling. In this embodiment, the atomic coordinates are selected from: (i) atomic coordinates determined by X-ray diffraction of a crystalline BCMA or crystalline BAFF-R; (ii) atomic coordinates selected from: (1) atomic coordinates represented in any one of Tables 13-33; (2) atomic coordinates that define a three dimensional structure having an average
30 root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional

structure represented by the atomic coordinates of (1); (3) atomic coordinates in any one of Tables 13-22 defining a portion of the BCMA, wherein the portion of the BCMA comprises sufficient structural information to perform step (b); and (4) atomic coordinates in any one of Tables 14-33 defining a portion of the BAFF-R, wherein the portion of the BAFF-R comprises sufficient structural information to perform step (b); and (iii) atomic coordinates defining the three dimensional structure of BCMA molecules or BAFF-R molecules arranged in a crystalline manner in a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217$.

Yet another embodiment of the invention relates to a method to construct a three dimensional model of BCMA, BAFF-R, TACI, or a homologue thereof, comprising: (a) obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R, the atomic coordinates being selected from any of those described above for BCMA or BAFF-R; and (b) performing computer modeling with the atomic coordinates of (a) and an amino acid sequence corresponding to BCMA, BAFF-R or TACI to construct a model of a three dimensional structure of the BCMA, BAFF-R or TACI, or homologue thereof.

Another embodiment of the invention relates to a crystal comprising a TALL-1 protein, wherein the crystal effectively diffracts X-rays for the determination of the atomic coordinates of the TALL-1 protein to a resolution of greater than 3.0\AA , and $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217$.

Brief Description of the Figures of the Invention

The patent or application file contains at least one drawing executed in color. Copies of this patent or patent application with color drawing will be provided by the Office upon request and payment of the necessary fee.

Fig. 1A is a ribbon diagram of the three-dimensional structure of sTALL-1 (residue 142-285); "elbow" and "flap" regions are unique for sTALL-1 and termed for their shapes; starting from the N-terminus, A (146-151) \rightarrow A' (158-160) \rightarrow A'' (163-165) \rightarrow A''' (168-174) \rightarrow B' (178-181) \rightarrow B (184-187) \rightarrow C (191-201) \rightarrow D (208-215) \rightarrow E (226-235) \rightarrow F (245-253) \rightarrow G (258-242) \rightarrow H (270-283).

Fig. 1B is a stereo view of superimposing sTALL-1 onto TNF α (PDB ID 1TNF); sTALL-1 is colored green; TNF α is colored yellow.

Fig. 1C is a ribbon representation of sTALL-1 trimer, looking down from the 3-axis fold that generates the trimer.

Fig. 1D is a stereo view of superimposing of sTALL-1 trimer and TNF α trimer, viewing from the orientation vertical to the 3-axis fold; sTALL1-1 is colored green, TNF α is colored gray.

Fig. 2A is a digitized image showing the relative position of an asymmetry unit in the cluster and overall structures of the cluster at different orientations; the 10 monomers (colored yellow) in the asymmetric unit, which generate the left 50 monomers (colored gray) through crystallographic symmetry ($P6_322$) to form the virus-like cluster with total of 60 monomers.

Fig. 2B is a digitized image showing the structure of virus-like cluster ($T=1$) looking down from the local 5-fold symmetry.

Fig. 2C is a digitized image showing the structure of virus-like cluster looking down from the 3-fold symmetry; all monomers are colored according to chains as default set in RIBBON.

Fig. 3A is a stereo view of the interactions between two sTALL-1 trimers; trimer 1 consists of monomers 1 (light gray), 2 (gray), and 3 (dark); trimer 2 contains monomers 1', 2', and 3'.

Fig. 3B shows the major interaction is involved in two monomers (monomer 1 and monomer 1'); two layered interactions are termed layer 1 (L1) and layer 2 (L2).

Fig. 3C shows residues and locations of layer 1.

Fig. 3D is a stereo view of the interactions of layer 1.

Fig. 3E shows residues and locations of layer 2.

Fig. 3F is a stereo view of the interactions of layer 2.

Fig. 3G shows residues and locations of the third layer interactions that involve three monomers; monomers 1 (yellow), 1' (gray), and 2' (dark).

Fig. 3H is a stereo view of the two hydrophobic cores formed by residues from three monomers (residues from monomer 1 are colored red, residues from monomer 1' are colored blue, residues from monomer 2' are colored dark).

Fig. 4A shows the electron microscopy view of sTALL-1 in solution after negative staining; black bar is 50nm long and clusters of sTALL-1 are around 20nm in diameter.

Fig. 4B is a surface presentation of sTALL-1 viewing from the similar orientation as figure 2B.

Fig. 4C is a surface presentation of sTALL-1 viewing from the similar orientation as figure 2C.

5 Fig. 5A shows a possible sub-cluster of four trimers (trimers 1, 2, 3, and 4) of sTALL-1 in the virus-like cluster.

Fig. 5B shows a possible sub-cluster of five trimers (trimers 1, 2, 3, 4, and 5) of sTALL-1.

10 Fig. 6A is an initial Of-Fc map of eBCMA with sTALL-1 at 2σ level; phases are calculated from sTALL-1 model (PDB ID, 1JH5); eBCMA is the final refined model; the map part is a representative of all eight binding receptors in the asymmetry unit, with most residues shown with their side-chains.

Fig. 6B is a ribbon diagram of the three-dimensional structure of eBCMA (residue 5-43 of SEQ ID NO:6); three disulfide bridges are also shown.

15 Fig. 6C is a ribbon diagram of the three-dimensional structure of eBAFF-R (residue 16-58 of SEQ ID NO:8); one disulfide bridge and two pseudo disulfide-like connections are also shown.

20 Fig. 7A shows the 60 monomers of sTALL-1 (colored green) and 60 monomers of eBAFF-R (molecules colored yellow are real from the complex structure, molecules colored blue are partially ordered, molecules colored red are missing in the complex due to crystal packing).

Fig. 7B shows a representation of Fig. 7A without sTALL-1.

Fig. 7C shows the 60 monomers of sTALL-1 and 60 monomers of eBCMA; all are colored according to secondary structure.

25 Fig. 7D shows a representation of Fig. 7C without sTALL-1.

Fig. 8A shows a superposition of eBCMA, eBAFF-R, and the C2 containing CRD from TNF-R1 (Naismith et al., 1998, *supra*).

30 Fig. 8B shows a structure based sequence alignment of CRD modules of BCMA (SEQ ID NO:11), BAFF-R (SEQ ID NO:12), TACI1 (SEQ ID NO:13), TACI2 (SEQ ID NO:14), Fn14 (SEQ ID NO:15), and TNF-R1 (SEQ ID NO:16); residues colored red are conserved disulfide bridges or pseudo disulfide bridges, which builds up module A1, D2, and

D0; residues colored yellow are not defined; residues colored blue are for the C2 module; residues colored green are putative residues involved in ligand recognition.

Fig. 9A shows the one to one mode interaction of eBCMA with sTALL-1.

Fig. 9B shows three eBCMA on the trimer sTALL-1.

5 Fig. 9C shows two trimers of eBCMA and sTALL-1 complex.

Fig. 9D shows the overall interactions between eBCMA and sTALL-1.

Fig. 9E shows the hydrophobic core 1 for the interaction between eBCMA and sTALL-1.

10 Fig. 9F shows the hydrophobic core 2 for the interaction between eBCMA and sTALL-1.

Fig. 9G shows salt bridges 1 and 2 for the interaction between eBCMA and sTALL-1.

Fig. 9H shows the overall interactions between eBAFF-R and sTALL-1.

Fig. 9I shows the hydrophobic core 1 for the interaction between eBAFF-R and sTALL-1.

15 Fig. 9J shows the hydrophobic core 2 for the interaction between eBAFF-R and sTALL-1.

Fig. 10A is a model of APRIL and its superposition on sTALL-1 in the presence of eBAFF-R; three sTALL-1 monomers are colored pink, three models of APRIL are colored gray, magenta, and blue respectively.

20 Fig. 10B shows a hypothetical overall interaction between eBAFF-R and APRIL.

Detailed Description of the Invention

The present invention relates to the determination of the three-dimensional structure of sTALL-1 and sTALL-1 in complex with the extracellular domains of its cognate receptors, BCMA and BAFF-R, and to the use of such structures to develop agonists and antagonists
25 and lead compounds for drug development in the area of therapeutic agents related to TALL-1 biological activity. The present invention specifically relates to various TALL-1 protein homologues (agonists and antagonists) that were designed using the structural information provided herein, as well as TALL-1 receptor antagonists that were designed in a similar
30 manner. The present invention also relates to agonist homologues of APRIL, and to the use of wild-type APRIL and such homologues in a method to inhibit the activity of TALL-1. The

present invention additionally relates to compositions comprising such homologues, agonists and antagonists, and to therapeutic methods of using such compounds and compositions. The invention further relates to crystalline complexes of sTALL-1 and sTALL-1 in complex with either BCMA or BAFF-R; to models of three-dimensional structures of such crystalline complexes and related structures, including models of the three dimensional structures of portions of the sTALL-1/BCMA complex or the sTALL-1/BAFF-R complex; to a method of drug design using any portion of such structures; to the design and/or identification of regulatory peptides derived from the knowledge of the three-dimensional structure of sTALL-1, the extracellular domains of BCMA, the extracellular domains of BAFF-R, and/or the complexes disclosed herein; to the compounds identified by drug design using such structures; and to the use of such compounds in therapeutic compositions and methods. These agents can be used to regulate B cell activity (e.g., B cell proliferation, B cell maturation, antibody production), autoimmunity, apoptosis, tumor cell survival, and other conditions affected by the activity of TALL-1, its receptors, and other TNF family members.

More particularly, the present inventors have determined the crystal structure of the functional soluble TALL-1 (sTALL-1) at 3.0 Å sTALL-1. The inventors have shown that the crystal structure of sTALL-1 forms a virus-like assembly with 200 Å diameter in the crystals, containing 60 sTALL-1 monomers. The cluster formation is mediated by a novel "flap" region of the sTALL-1 monomer. The virus-like assembly was also detected in solution using gel-filtration and electron microscopy. Deletion of the "flap" region disrupted the ability of TALL-1 monomers to form the virus-like assembly. Moreover, the mutant sTALL-1 bound its receptor, but could not activate NF-κB and did not stimulate B lymphocyte proliferation. Finally, the inventors found that the virus-like cluster of sTALL-1 exists in physiological conditions. Details of the structure of TALL-1 are discussed below.

In addition, the present inventors have determined the crystal structures of sTALL-1 complexed with the extracellular domains of BCMA and BAFF-R at 2.6Å and 3.0Å, respectively. The single cysteine rich domain (CRD) of BCMA and BAFF-R both have a saddle-like architecture, which sits on the horseback-like groove formed by four coil regions on each individual sTALL-1 monomer. Two novel structural modules D2 and D0 were revealed from these structures. Details of the structure of sTALL-1 in complex with its cognate receptors are also discussed below.

Using the information provided herein regarding the structure of TALL-1 and its receptors, one can design agonists and antagonists of the both TALL-1 and the receptors. The present inventors have identified the residues of TALL-1 that are important for trimer formation, for the interaction between trimers of TALL-1, and for binding of TALL-1 to both BCMA and BAFF-R. Additionally, the inventors have determined residues of BCMA and BAFF-R that are important for binding to TALL-1, and this information is predictive of TALL-1-binding residues of the third known receptor for TALL-1, TACI.

Finally, from sequence alignments, the truncated version of sTALL-1 is similar to APRIL (TALL-2), the closely related family member of TALL-1. Moreover, all residues that take part in the trimer-trimer interactions are not conserved between TALL-1 and TALL-2. It seems impossible for APRIL to form the virus-like cluster. It is suggested that APRIL may act as a decoy ligand *in vivo*. The inability of APRIL to bind to BAFF-R indicates structural diversity between TALL-1 and APRIL (Schiemann et al., (2001) *Science* 293:2111-2114). The resolution of the TALL-1 structure described herein has allowed the present inventors to model the related protein, APRIL, and to propose novel agonists of APRIL, as well as a novel function for APRIL, as a decoy ligand for TALL-1.

As demonstrated by multiple laboratories, administration of sTALL-1 can cause autoimmune like lupus in mice (Gross et al., (2000) *Nature*, 404:995-999; Mackay et al., (1999) *J Exp Med.* 190:1697-710; Khare et al., (2000) *Proc Natl Acad Sci USA.* 97:3370-5). The present inventors reason that a non-functional mutation of sTALL-1, which still has similar binding affinity to its receptors competing with native sTALL-1, could serve as a therapeutic candidate for treating autoimmune diseases. The truncated version of sTALL-1 lacks the ability to form clusters, is defective in NF κ B activation function, but still binds to its cognate receptor, making it a possible candidate for this purpose. Therefore, the present inventors' discoveries have applications for designing novel TALL-1 antagonists (and agonists) and novel TALL-1 receptor antagonists (and agonists) for use in therapeutics to regulate B cell activity (e.g., B cell proliferation, B cell maturation, antibody production), autoimmunity, apoptosis, tumor cell survival, and other conditions affected by the activity of TALL-1, its receptors, and other TNF family members.

According to the present invention, general reference to TALL-1 (e.g., BAFF, THANK, BlyS or zTNF4) refers to a tumor necrosis factor (TNF)/tumor necrosis factor

receptor (TNFR) family member which has been characterized as playing a role in B cell development and maturation (Shu et al., 1999, *J. Leukocyte Biology* 65:680-683; Schneider et al., 1999, *J Exp Med.* 189:1747-56; Moore et al., 1999, *Science* 285:260-263; Mukhopadhyay et al., 1999, *supra*; Shu et al., 2000, *Pro. Natl. Acad. Sci. USA* 97:9156-9161; Gross et al., 2000, *Nature* 404:995-999; Thompson et al., 2000, *J Exp Med* 192:129-35; Marsters et al., 2000, *Curr Biol.* 10:785-8; Xia et al., 2000, *J Exp Med.* 192:137-43; Yan et al., 2000, *Nat. Immunol.* 1:37-41; Thompson et al., 2001, *Science* 293: 2108-2111). The amino acid sequence of TALL-1 is represented herein by SEQ ID NO:2. SEQ ID NO:2 (encoded by the nucleic acid sequence SEQ ID NO:1) represents the full-length TALL-1 protein sequence. Amino acid positions for TALL-1 described herein are made with reference to SEQ ID NO:2, unless otherwise noted. The amino acid sequence of soluble TALL-1 (sTALL-1) is consists of positions 134 to 285 of SEQ ID NO:2. In general, reference to a TALL-1 protein can include both the full-length TALL-1 represented by SEQ ID NO:2 and the soluble TALL-1 represented by positions 134-285 of SEQ ID NO:2. The crystal structure of the sTALL-1 protein described herein comprises amino acid positions 134-285 of SEQ ID NO:2. The TALL-1 protein used for crystallization included an N-terminal His₆ tag, facilitating isolation and purification using nickel-chelating affinity chromatography.

The present inventors have determined that the structure of sTALL-1 consists of two layered antiparallel β strands that form a typical jellyroll-like β sandwich, as with other members of the TNF ligand family (Jones et al., (1989) *Nature* 338:225-228; Eck et al., (1989) *J. Biol. Chem.*, 264:17595-17605; Eck et al., (1992) *J. Biol. Chem.* 267:2119-2122; Karpusas et al., (1995) *Structure* 3:1031-1039; Cha et al., 1999, *Immunity* 11:253-261). Compared to known structures of other family members, the overall structure of sTALL-1 is shorter along the 3-fold axis that generates the trimers (Fig. 1B). Two unique features of TALL-1 are termed "elbow" and "flap" regions (Fig. 1A). The "elbow" region contains a short β hair-pin labeled A" and A". The "flap" region is unique to sTALL-1 based on results of sequence alignments and structural comparisons (Figs. 1B, 1D). The unique "flap" region of sTALL-1 mediates trimer-trimer interactions that lead to a remarkable virus-like assembly of the sTALL-1 trimers. There are 10 sTALL-1 monomers in the asymmetric unit with a space group of $P6_322$ (Fig. 2A). The 10 monomers interact to form virus-like clusters

containing 60 sTALL-1 monomers (20 trimers) (Figs. 2B, 2C). The trimer-trimer interactions are extensive. They not only include hydrogen bond net works and salt bridges, but also hydrophobic contacts. Residues involved in trimer-trimer interactions are not only from the monomer that contributes the "flap" region but also the neighboring monomer as well (Figs 3A and 3B). The fine details of the structure of TALL-1 are described in Example 1. The present inventors additionally showed that the virus-like cluster assembly could be visualized by electron microscopy, that the cluster assembly exists in solution as physiological pH, and that the "flap region" (and by extension the ability to form clusters) was essential for the proper function of sTALL-1 *in vivo* (see Example 2).

According to the present invention, general reference to a receptor for TALL-1 or a "TALL-1 receptor" generally refers to any of the cognate receptors for TALL-1, including the receptors known as BCMA (B cell maturation factor), BAFF-R (also called BR3), and TACI (Shu et al., 2000, *Pro. Natl. Acad. Sci. USA.* 97:9156-9161; Gross et al., 200, *Nature* 404:995-999; Thompson et al., 2000, *J. Exp Med.* 192:129-35; Marsters et al., 2000, *Curr Biol.* 10:785-8; Xia et al., 2000, *J Exp Med.* 192:137-43; Yan et al., 2000, *Nat. Immunol.* 1:37-41; Thompson et al., 2001, *Science* 293:2108-2111; Yan et al., 2001, *Curr Biol.* 11:1547-1552; each of which is incorporated herein by reference in its entirety).

The amino acid sequence of BCMA is represented herein by SEQ ID NO:6. SEQ ID NO:6 (encoded by the nucleic acid sequence SEQ ID NO:5) represents the full-length BCMA protein sequence. Amino acid positions for BCMA described herein are made with reference to SEQ ID NO:6, unless otherwise noted. A soluble BCMA can include positions 1-62 of SEQ ID NO:6, or a smaller fragment within positions 1-62 of SEQ ID NO:6. The crystal structure of the eBCMA protein (extracellular domain of BCMA) described herein was produced using amino acid residues 1-52 of SEQ ID NO:6; the residues ordered in the structure model of eBCMA described herein comprises residues 5-43 of SEQ ID NO:6 (Fig. 6B). The amino acid sequence of BAFF-R is represented herein by SEQ ID NO:8. SEQ ID NO:8 (encoded by the nucleic acid sequence SEQ ID NO:7) represents the full-length BAFF-R protein sequence. Amino acid positions for BAFF-R described herein are made with reference to SEQ ID NO:8, unless otherwise noted. The crystal structure of the eBAFF-R protein (extracellular domain of BAFF-R) described herein was produced using amino acid residues 1-62 of SEQ ID NO:8; the residues ordered in the structure model of eBAFF-R

described herein comprises residues 16-58 of SEQ ID NO:8 (Fig. 6C). The eBCMA and the eBAFF-R used for crystallization included a GST tag, facilitating isolation and purification using affinity chromatography.

The present inventors have determined the structure of the sTALL-1 (described above) in complex with each of eBCMA and eBAFF-R. The space group of the TALL-1 crystals remained $P6_322$ with the same cell dimensions with or without binding of the receptors. There were two virus-like clusters of TALL-1 in one unit cell, and each cluster had 60 copies of sTALL-1, 42 fully occupied eBCMA or eBAFF-R, and 6 partial copies of eBCMA or eBAFF-R. There were 12 copies of sTALL-1 free of receptors due to crystal packing. All receptors were located on the outer-extreme shell, which expands the ball-like shell another ~ 20 Å in each direction. The overall arrangement of the receptors on the shell resembled a sunflower with receptors as flower petals and sTALL-1 as a seed bed (Fig. 7). The interactions between sTALL-1 and eBAFF-R are similar to those between sTALL-1 and eBCMA, although details are slightly different. The interaction modes of the eBCMA and eBAFF-R with sTALL-1 are dramatically different from those found in the other TNF family members, containing at least two CRDs that bind to the cleft regions formed by two ligands. For the interactions described here, one saddle-like receptor mostly makes a one to one interaction with its ligand at the extreme end of the ligand (Figs. 9A-9C). The difference exists not only in the CRD structure but also in the binding locations and modes.

The sequence homology between eBCMA and eTACI (extracellular domain of TACI) is obvious. This is not true for eBCMA and eBAFF-R or for eTACI and eBAFF-R. The structures of eBCMA and eBAFF-R allowed the present inventors to perform a structural based sequence alignment of eBCMA, eBAFF, and eTACI. They found that a strong pattern of similarity emerges (Fig. 8A and 8B), and thus it can be predicted that TACI will bind to TALL-1 in a manner similar to that of BCMA and BAFF-R described herein. Details regarding the structure of eBCMA and eBAFF-R and the interaction between these receptors and TALL-1 are described in Example 6.

The present inventors' results are not consistent with two published results (Schneider et al., 1999, *J Exp Med* 189:1747-56; and Kanakaraj et al., 2001, *Cytokine* 13:25-31), both of which claimed that sTALL-1/BAFF existed only as trimers. Moreover, two publications of TALL-1 structure published subsequent to the filing of the priority document for this

application did not report the assembly of TALL-1 monomers into the virus-like cluster described herein (Oren et al., Feb 25, 2002, *Nat. Struct. Biol.* 9(4):288-292; Karpusas et al., Feb 1 2002, *J. Mol. Biol.* 315:1145-1154).

However, four lines of evidence support the present inventors' belief that the crystal structure described herein reflects the actual interactions of the complexes in solution and *in vivo*. First, co-expression of sTALL-1 with eBCMA or eBAFF-R generates the virus-like cluster in solution as judged by gel-filtration column and SDS-phage analysis at a ratio of 1:1 (sTALL-1: eBCMA or eBAFF-R). Different salt concentrations (from 100mM to 1M NaCl) produce the same elution profile, in which complexes of sTALL-1 with eBCMA or eBAFF-R elute at a void volume on superdex-200. Thus, binding between ligand and receptors is stable and insensitive to salt concentrations. Purified samples of the preformed complexes were subjected to crystallization trials. Crystals of both complexes have been obtained, however neither of them diffracted. The results further confirmed that the "flap" region of TALL-1 which is involved in clustering, is not part of the receptor binding site. Furthermore, these results suggest that the pre-binding of receptors to sTALL-1 disrupts the original molecular packing in the sTALL-1 crystals and that the receptors are located on the surface of the sTALL-1 cluster. Second, in the receptor soaked sTALL-1 crystals, all seven fully occupied receptors and one partial receptor have equivalent binding sites on sTALL-1 in the asymmetry unit, so the binding is highly specific. Third, eBCMA and eBAFF-R have a similar binding mode and occupy the same site on sTALL-1. Fourth, each of the three C-termini of eBCMA and eBAFF-R on the sTALL-1 trimer point to the same direction, the putative membrane surface for trimerization (Fig. 7C and 7D). Therefore, without being bound by theory, the present inventors believe that the interactions revealed from the complex structures represent the actual interactions between TALL-1/BCMA and TALL-1/BAFF-R *in vivo*.

Finally, the present inventors have modeled APRIL based on the sTALL-1 structure, benefitting from the high primary sequence homology between TALL-1 and APRIL. According to the present invention, general reference to APRIL refers to a tumor necrosis factor (TNF)/tumor necrosis factor receptor (TNFR) family member which is the closest family member to TALL-1, and is represented herein by SEQ ID NO:4 (encoded by the nucleic acid sequence SEQ ID NO:3). APRIL has low abundance in normal tissues, but is

present at high level in transformed cell lines and in variety of human cancers (Hahne et al., 1998, *J. Exp. Med.* 188:1185-1190). More recent data show that BAFF-R does not bind APRIL (Thompson et al., 2001, *Science* 293:2108-2111; Yan et al., 2001, *Curr Biol.* 11:1547-1552), suggesting that APRIL is dispensable for B cell maturation (Thompson et al., 5 2001, *supra*; Yan et al., 2001, *supra*; Schneider et al., 2001, *J. Exp. Med.* 194:1691-1697), although it binds to BCMA and TACI with an affinity similar to sTALL-1 (Yu et al. 2000, *Nature immunol.* 1:252-256). Nevertheless, APRIL-deficient mice die in utero (Mackay et al., 2002, *TRENDS in Immunology* 23:113-115), leaving the role of APRIL *in vivo* at the time of this invention a mystery.

10 The final built model of APRIL was imported to the minimization program in CNS (Brunger et al., 1998, *Acta Cryst* D54:905-921, and the output coordinates were superimposed on the sTALL-1 structure (Fig. 10A). All residues from eBAFF-R that are involved in the interactions between the eBAFF-R and sTALL-1 are displayed (Fig. 10B). All equivalent residues in APRIL, which are close to the receptor binding surface in sTALL-1 15 are also shown (Fig. 10B). To the present inventors' surprise, the interactions were extremely similar to those found in the complexes of eBCMA or eBAFF-R with sTALL-1. The most obvious difference between sTALL-1 and APRIL is in the "flap" region (8 residues; 217-224 of SEQ ID NO:2) of sTALL-1, which is missing in APRIL (Shu et al., 1999, *J. Leukocyte Biology* 65:680-683). The present inventors have reported a mutated 20 version of sTALL-1 with 8 residues of the "flap" region replaced by two glycine residues, and this mutant was not functional in transfection assays or in the B-cell stimulation assays, but had a binding affinity to its receptors similar to that of the native sTALL-1. The present inventors have determined the structure of this mutated sTALL-1 at 1.7 Å resolution by MIR, and it is almost identical to the sTALL-1 except for the missing flap (data not shown). 25 Moreover, this mutated sTALL-1 is a close model of APRIL. Therefore, without being bound by theory, the present inventors believe that APRIL may be serving as a decoy ligand, reducing the opportunity for sTALL-1 to bind to the same receptor. This role is similar to the decoy death receptors, which are essential for cells to survive (Cha et al., 1999, *Immunity* 11:253-261; Mongkolsapaya et al., 1999, *Nat. Struct. Biol.* 6:1048-1053; Hymowitz et al., 30 1999, *Mol. Cell* 4:563-571). As shown in Example 6, although APRIL does not bind to BAFF-R under physiological conditions (pH7.5), the present inventors have produced at least

two homologues of BAFF-R that can bind to APRIL under physiological conditions. Also, the present inventors have demonstrated that APRIL can form heterotrimers with sTALL-1 under physiological conditions. These results and the implications therefore are discussed in detail below.

5 As discussed above, various details of the structure of TALL-1, of the TALL-1 receptors BCMA and BAFF-R, and of the interactions between TALL-1 monomers and between TALL-1 and its receptors are described herein and particularly in the Examples section. This information can now be used to design novel agonists and antagonists of TALL-1 and its cognate receptors, embodiments of which are described in detail below.

10 Accordingly, one embodiment of the present invention relates to a variety of TALL-1 homologues and particularly, TALL-1 agonist and TALL-1 antagonist proteins, that are designed using the structural information provided herein. The following discussion is made with reference to TALL-1 proteins, including homologues thereof, but it is to be understood, however, that the general definitions of terms and methods are intended to apply to the
15 discussion of an isolated TALL-1 receptor and homologues thereof, as well as to discussion of APRIL and homologues thereof, unless otherwise modified within the specific discussion of the TALL-1 receptor or APRIL.

An isolated protein (e.g., an isolated TALL-1 protein), according to the present invention, is a protein that has been removed from its natural milieu (i.e., that has been
20 subject to human manipulation) and can include purified proteins, partially purified proteins, recombinantly produced proteins, and synthetically produced proteins, for example. As such, "isolated" does not reflect the extent to which the protein has been purified. Preferably, an isolated protein, and particularly, an isolated TALL-1 (including fragments and homologues thereof), is produced recombinantly. The terms "fragment", "segment" and "portion" can be
25 used interchangeably herein with regard to referencing a part of a protein. It will be appreciated that, as a result of the determination of the tertiary structure of biologically active portions of TALL-1 and the extracellular domains of two of its receptors herein, various portions and residues of TALL-1 and its receptors will now be considered to be particularly valuable for mutational analyses and various biological assays, as well as for the
30 development of therapeutic proteins and compounds or lead compounds for drug design, and also for computer-assisted drug design methods, as discussed herein. Such portions of

TALL-1 and its receptors and methods of using such portions are explicitly contemplated to be part of the present invention.

According to the present invention, general reference to TALL-1 is reference to a protein that typically contains any biologically active portion of a native or wild-type TALL-1 protein (e.g., a portion that can exhibit at least one biological activity associated with native (wild-type) TALL-1 or a portion that at least binds to a given receptor), and includes full-length TALL-1, soluble proteins, biologically active fragments of TALL-1, TALL-1 fusion proteins, or any homologue of a naturally occurring TALL-1, as described in detail below. Similarly, general reference to a TALL-1 receptor is reference to a protein that typically contains any biologically active portion of a native or wild-type TALL-1 receptor (e.g., a portion that can exhibit at least one biological activity associated with native (wild-type) TALL-1 receptor or a portion that at least binds to a given ligand, such as TALL-1), and includes full-length TALL-1 receptor, soluble receptors, biologically active fragments of TALL-1 receptors, TALL-1 receptor fusion proteins, or any homologue of a naturally occurring TALL-1 receptor, as described in detail below. General reference to APRIL herein is a reference to a protein that typically contains any biologically active portion of a native or wild-type APRIL protein (e.g., a portion that can exhibit at least one biological activity associated with native (wild-type) APRIL or a portion that at least binds to a given receptor), and includes full-length APRIL, soluble proteins, biologically active fragments of APRIL, APRIL fusion proteins, or any homologue of a naturally occurring APRIL (including both agonists and antagonists), as described in detail below.

Reference herein to a protein from a specific organism, such as a "human TALL-1", by way of example, refers to a TALL-1 protein from a human or to a TALL-1 protein that has been otherwise produced from the knowledge of the primary structure (e.g., sequence) and/or the tertiary structure of a naturally occurring TALL-1 protein from *Homo sapiens*. In other words, a human TALL-1 protein includes any TALL-1 protein that has the structure and function of a naturally occurring TALL-1 from *Homo sapiens* or that has a structure and function that is sufficiently similar to a human TALL-1 such that the TALL-1 protein is a homologue of a naturally occurring TALL-1 from *Homo sapiens*. As such, a human TALL-1 protein, by way of example, can include purified, partially purified, recombinant, mutated/modified and synthetic proteins.

A homologue of any protein described herein includes proteins which differ from a naturally occurring protein in that at least one or a few, but not limited to one or a few, amino acids have been deleted (e.g., a truncated version of the protein, such as a peptide or fragment), inserted, inverted, substituted and/or derivatized (e.g., by glycosylation, phosphorylation, acetylation, myristoylation, prenylation, palmitation, amidation and/or addition of glycosylphosphatidyl inositol). In other words, a homologue of a protein according to the invention includes proteins that have been mutated or modified, as compared to the wild-type protein. According to the present invention, the terms "modification" and "mutation" can be used interchangeably, particularly with regard to the modifications/mutations to the amino acid sequences of TALL-1, APRIL or TALL-1 receptors (or nucleic acid sequences) described herein. A homologue can have either enhanced, decreased, or substantially similar properties (including combinations thereof, when different properties are assessed) as compared to the naturally occurring protein or peptide. A homologue can include an agonist of a protein or an antagonist of a protein.

Homologues can be produced using techniques known in the art for the production of proteins including, but not limited to, direct modifications to the isolated, naturally occurring protein, direct protein synthesis, or modifications to the nucleic acid sequence encoding the protein using, for example, classic or recombinant DNA techniques to effect random or targeted mutagenesis.

Modification of proteins described herein typically result in homologues that have agonistic and/or antagonistic biological activities as compared to the naturally occurring (wild-type) protein. In general, the biological activity or biological action of a protein refers to any function(s) exhibited or performed by the protein that is ascribed to the naturally occurring form of the protein as measured or observed *in vivo* (i.e., in the natural physiological environment of the protein) or *in vitro* (i.e., under laboratory conditions). Modifications of a protein, such as in a homologue or mimetic (discussed below), may result in proteins having the substantially the same biological activity as the naturally occurring protein, or in proteins having decreased or increased biological activity as compared to the naturally occurring protein. Modifications which result in a decrease in protein expression or a decrease in the activity of the protein, can be referred to as inactivation (complete or partial), down-regulation, or decreased action of a protein. Similarly, modifications which

result in an increase in protein expression or an increase in the activity of the protein, can be referred to as amplification, overproduction, activation, enhancement, up-regulation or increased action of a protein.

According to the present invention, an isolated protein described herein, including a
5 biologically active homologue (agonist or antagonist) or fragment thereof, has at least one characteristic biological activity of the wild-type, or naturally occurring protein (which can vary depending on whether the homologue or fragment is an agonist, antagonist, or mimic of the wild-type protein). For example, the biological activity of TALL-1 can include, but is not limited to, binding to at least one TALL-1 receptor (e.g., BCMA, BAFF-R, TACI);
10 activation of at least one TALL-1 receptor; formation of trimers with other TALL-1 monomers, formation of viral-like clusters among TALL-1 trimers, an ability to costimulate B lymphocyte proliferation; an ability to costimulate B lymphocyte activation; and/or an ability to support B lymphocyte survival and development. Biological activity of a TALL-1 receptor can include, but is not limited to: an ability to bind to a ligand, including TALL-1
15 or APRIL; receptor translocation within a cell upon ligand binding; NF κ B activation; TRAF5, TRAF6, NIK, IKK α and/or IKK β activation; costimulation of B cell proliferation; costimulation of B cell activation; and enhancement of B cell survival. Biological activity of APRIL can include, but is not limited to: binding to at least one APRIL receptor (e.g., BCMA, TACI); regulation of B cell survival, development or maturation. Biological
20 activities of TALL-1, APRIL and TALL-1/APRIL receptors are known in the art and are described in: Shu et al., 1999, *J. Leukocyte Biology* 65:680-683; Schneider et al., 1999, *J Exp Med.* 189:1747-56; Moore et al., 1999, *Science* 285:260-263; Mukhopadhyay et al., 1999, *supra*; Shu et al., 2000, *Pro. Natl. Acad. Sci. USA* 97:9156-9161; Gross et al., 2000, *Nature* 404:995-999; Thompson et al., 2000, *J Exp Med* 192:129-35; Marsters et al., 2000,
25 *Curr Biol.* 10:785-8; Xia et al., 2000, *J Exp Med.* 192:137-43; Yan et al., 2000, *Nat. Immunol.* 1:37-41; Yu et al., 2000, *Nat Immunol.* 1:252-6; Thompson et al., 2001, *Science* 293: 2108-2111; and Yan et al., 2001, *Curr Biol.* 11(19):1547-52; each of which is incorporated herein by reference in its entirety.

Methods of detecting and measuring such biological activity, including measuring
30 agonist or antagonist activity, include, but are not limited to measurement of transcription of the protein; measurement of translation of the protein; measurement of secretion of soluble

forms of the protein (TALL-1 and APRIL); measurement of binding of the protein to its receptor (TALL-1 and APRIL); measurement of binding of the protein to its ligand (BCMA, BAFF-R, TACT); measurement of B cell proliferation; measurement of B cell activation; measurement of B lymphocyte cytokine production; measurement of NF κ B activation;
5 measurement of TRAF5, TRAF6, NIK, IKK α or IKK β activation; measurement of immunoglobulin maturation; measurement of immunoglobulin production and secretion; measurement of calcium mobilization; or measurement of phosphorylation of signal transduction proteins. It is noted that a homologue of a protein according to the present invention is not required to have all of the biological activities of the wild-type protein. For
10 example, a TALL-1 homologue may bind to a TALL-1 receptor, but may not be able to activate the receptor. Various homologues are useful as agonists or antagonists of the wild-type protein and in addition, some homologues are useful in diagnostic assays, as lead compounds for drug design, or in screening assays, for example, or for other purposes such as antibody production.

15 In general, methods to measure protein expression levels include, but are not limited to: western blotting, immunocytochemistry, flow cytometry or other immunologic-based assays; assays based on a property of the protein including but not limited to DNA binding, ligand binding, or interaction with other protein partners. Binding assays are also well known in the art. For example, a BIAcore machine can be used to determine the binding
20 constant of a complex between two proteins. The dissociation constant for the complex can be determined by monitoring changes in the refractive index with respect to time as buffer is passed over the chip (O'Shannessy et al. Anal. Biochem. 212:457-468 (1993); Schuster et al., Nature 365:343-347 (1993)). Other suitable assays for measuring the binding of one protein to another include, for example, immunoassays such as enzyme linked
25 immunoabsorbent assays (ELISA) and radioimmunoassays (RIA), or determination of binding by monitoring the change in the spectroscopic or optical properties of the proteins through fluorescence, UV absorption, circular dichroism, or nuclear magnetic resonance (NMR).

Measurement of expression of transcripts can be performed by any of a variety of
30 known methods in the art. For RNA expression, methods include but are not limited to, extraction of cellular mRNA and northern blotting using labeled probes that hybridize to

transcripts encoding all or part of the mRNA encoding the protein of interest; amplification of mRNA using sequence-specific primers and reverse transcriptase-polymerase chain reaction (RT-PCR), followed by quantitative detection of the product by any of a variety of means; extraction of total RNA from the cells, which is then labeled and used to probe
5 cDNAs. The term "quantifying" or "quantitating" when used in the context of quantifying transcription levels of a gene can refer to absolute or to relative quantification. Absolute quantification may be accomplished by inclusion of known concentration(s) of one or more target nucleic acids and referencing the hybridization intensity of unknowns with the known target nucleic acids (e.g. through generation of a standard curve). Alternatively, relative
10 quantification can be accomplished by comparison of hybridization signals between two or more genes, or between two or more treatments to quantify the changes in hybridization intensity and, by implication, transcription level.

Included in the invention are both agonists and antagonists of proteins described herein. As used herein, reference to an agonist, as in a "TALL-1 agonist" or "APRIL agonist"
15 refers to any compound that is characterized by the ability to agonize (e.g., stimulate, induce, increase, enhance, or mimic) the biological activity of the naturally occurring protein (TALL-1 or APRIL, respectively) as described herein (e.g., by interaction/binding with and/or activation of a receptor for the naturally occurring protein). More particularly, an agonist as set forth above can include any compound that selectively binds to and/or activates or
20 increases the activation of a TALL-1 receptor or APRIL receptor, respectively, or otherwise mimics or enhances the activity of the natural ligand, TALL-1 or APRIL, respectively. Similarly, reference to a "TALL-1 receptor agonist" or "APRIL receptor agonist" refers to any compound that is characterized by its ability to agonize (e.g., stimulate, induce, increase, enhance) the biological activity of the naturally occurring receptor as described herein (e.g.,
25 by interaction/binding with ligands of the receptor and mimicking or enhancing the biological activity of the receptor). Agonists can include, but are not limited to, a protein, a peptide, a nucleic acid, or any product of drug/compound/peptide design or selection and includes any homologue of the protein, binding protein (e.g., an antibody), agent, or any suitable product of drug/compound/peptide design or selection which is characterized by its ability to agonize
30 (e.g., stimulate, induce, increase, enhance) the biological activity of the naturally occurring protein (e.g., TALL-1) in a manner similar to the natural agonist (e.g., TALL-1). Agonists

of TALL-1, and TALL-1 receptors of the present invention can be useful in methods for increasing B cell development, B cell proliferation and/or B cell survival. Such agonists might be useful, for example, in conditions or diseases where B lymphocyte deficiency or hypoproliferation is problematic. Interestingly, the present inventors believe that APRIL agonists (compounds that agonize the activity of APRIL) can effectively serve as TALL-1 antagonists (e.g., they may antagonize the activity of TALL-1).

The phrase, "antagonist", as in a "TALL-1 antagonist" or "APRIL antagonist" refers to any compound which inhibits (e.g., antagonizes, reduces, decreases, blocks, reverses, or alters) the effect of a TALL-1 agonist or an APRIL agonist, respectively, as described above. More particularly, a TALL-1 antagonist or APRIL antagonist is capable of associating with a receptor (e.g., a TALL-1 receptor or an APRIL receptor, respectively), or otherwise acts in a manner relative to TALL-1 or APRIL activity, respectively, such that the biological activity of the receptor or of the natural agonist, is decreased in a manner that is antagonistic (e.g., against, a reversal of, contrary to) to the natural action of natural agonist. For example, an antagonist of TALL-1 could competitively inhibit the interaction between a natural TALL-1 and its receptor, and/or could induce a different effect on the receptor as compared to the effect induced by TALL-1. Similarly, a TALL-1 receptor antagonist or an APRIL receptor antagonist is capable of mimicking the structure of the natural receptor, and/or associating with TALL-1 or APRIL, respectively, in a manner that is antagonistic (e.g., against, a reversal of, contrary to) to the natural action of the receptor or the receptor upon binding to the natural ligand. Such antagonists can include, but are not limited to, a protein, peptide, a nucleic acid (including ribozymes and antisense) or product of drug/compound/peptide design or selection that provides the antagonistic effect. Antagonists of TALL-1 or TALL-1 receptor antagonists can be useful in methods for decreasing B cell development, B cell proliferation and/or B cell survival, such as in conditions or diseases where B cell hyperproliferation, or inappropriate B cell development or survival (e.g., autoimmune disease) is problematic.

Proteins of the present invention, including homologues, are preferably retrieved, obtained, and/or used in "substantially pure" form. As used herein, "substantially pure" refers to a purity that allows for the effective use of the protein *in vitro*, *ex vivo* or *in vivo* according to the present invention. For a protein to be useful in an *in vitro*, *ex vivo* or *in vivo* method according to the present invention, it is substantially free of contaminants, other proteins

and/or chemicals that might interfere or that would interfere with its use in a method disclosed by the present invention, or that at least would be undesirable for inclusion with the protein when it is used in a method disclosed by the present invention. For example, for a TALL-1 protein, such methods can include crystallization of the protein or use of all or a portion of the protein for mutational analysis, for antibody production, for agonist/antagonist identification assays, and all other methods disclosed herein. For a TALL-1 antagonist protein, such methods include use of the antagonist in a therapeutic composition or in a screening assay. Preferably, a "substantially pure" protein, as referenced herein, is a protein that can be produced by any method (i.e., by direct purification from a natural source, recombinantly, or synthetically), and that has been purified from other protein components such that the protein comprises at least about 80% weight/weight of the total protein in a given composition (e.g., the protein is about 80% of the protein in a solution/composition/buffer), and more preferably, at least about 85%, and more preferably at least about 90%, and more preferably at least about 91%, and more preferably at least about 92%, and more preferably at least about 93%, and more preferably at least about 94%, and more preferably at least about 95%, and more preferably at least about 96%, and more preferably at least about 97%, and more preferably at least about 98%, and more preferably at least about 99%, weight/weight of the total protein in a given composition.

A variety of homologues of TALL-1, APRIL and TALL-1 receptors are described herein. Although specific differences between a homologue and the wild-type protein are described below, in general, the homologue may have other modifications that do not necessarily effect the structure or biological activity of the homologue, but which cause it to have a different linear sequence as compared to the wild-type sequence. For example, a homologue having specified substantive modifications may also have multiple conservative amino acid substitutions so that the overall sequence identity between the wild-type protein and the homologue is less than if just the specified substantive modifications were made. Conservative substitutions typically include substitutions within the following groups: glycine and alanine; valine, isoleucine and leucine; aspartic acid, glutamic acid, asparagine, and glutamine; serine and threonine; lysine and arginine; and phenylalanine and tyrosine. Substitutions may also be made on the basis of conserved hydrophobicity or hydrophilicity (Kyte and Doolittle, *J. Mol. Biol.* (1982) 157: 105-132), or on the basis of the ability to

assume similar polypeptide secondary structure (Chou and Fasman, *Adv. Enzymol.* (1978) 47: 45-148, 1978). Therefore, in general, a homologue according to the present invention (e.g., a TALL-1 homologue, an APRIL homologue, or a TALL-1 receptor homologue) has an amino acid sequence that is at least about 50% identical to the amino acid sequence of the naturally occurring, or wild-type protein, (e.g., for TALL-1, the wild-type protein is represented herein as SEQ ID NO:2), and in another aspect at least about 55%, and in another aspect at least about 60%, and in another aspect at least about 65%, and in another aspect at least about 75%, and in another aspect at least about 75%, and in another aspect at least about 80%, and in another aspect at least about 85%, and in another aspect at least about 90%, and in another aspect at least about 95% identical to the amino acid sequence of the naturally occurring protein.

As used herein, unless otherwise specified, reference to a percent (%) identity refers to an evaluation of homology which is performed using: (1) a BLAST 2.0 Basic BLAST homology search using *blastp* for amino acid searches, *blastn* for nucleic acid searches, and *blastX* for nucleic acid searches and searches of translated amino acids in all 6 open reading frames, all with standard default parameters, wherein the query sequence is filtered for low complexity regions by default (described in Altschul, S.F., Madden, T.L., Schääffer, A.A., Zhang, J., Zhang, Z., Miller, W. & Lipman, D.J. (1997) "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs." *Nucleic Acids Res.* 25:3389-3402, incorporated herein by reference in its entirety); (2) a BLAST 2 alignment (using the parameters described below); (3) and/or PSI-BLAST with the standard default parameters (Position-Specific Iterated BLAST). It is noted that due to some differences in the standard parameters between BLAST 2.0 Basic BLAST and BLAST 2, two specific sequences might be recognized as having significant homology using the BLAST 2 program, whereas a search performed in BLAST 2.0 Basic BLAST using one of the sequences as the query sequence may not identify the second sequence in the top matches. In addition, PSI-BLAST provides an automated, easy-to-use version of a "profile" search, which is a sensitive way to look for sequence homologues. The program first performs a gapped BLAST database search. The PSI-BLAST program uses the information from any significant alignments returned to construct a position-specific score matrix, which replaces the query sequence for the next

round of database searching. Therefore, it is to be understood that percent identity can be determined by using any one of these programs.

Two specific sequences can be aligned to one another using BLAST 2 sequence as described in Tatusova and Madden, (1999), "Blast 2 sequences - a new tool for comparing protein and nucleotide sequences", *FEMS Microbiol Lett.* 174:247-250, incorporated herein
5 by reference in its entirety. BLAST 2 sequence alignment is performed in blastp or blastn using the BLAST 2.0 algorithm to perform a Gapped BLAST search (BLAST 2.0) between the two sequences allowing for the introduction of gaps (deletions and insertions) in the resulting alignment. For purposes of clarity herein, a BLAST 2 sequence alignment is
10 performed using the standard default parameters as follows.

For blastn, using 0 BLOSUM62 matrix:

Reward for match = 1

Penalty for mismatch = -2

Open gap (5) and extension gap (2) penalties

15 gap x_dropoff (50) expect (10) word size (11) filter (on)

For blastp, using 0 BLOSUM62 matrix:

Open gap (11) and extension gap (1) penalties

gap x_dropoff (50) expect (10) word size (3) filter (on).

20 In one aspect, a homologue of a protein described herein can also include proteins having an amino acid sequence comprising at least 25 contiguous amino acid residues of the wild-type sequence (i.e., 25 contiguous amino acid residues having 100% identity with 25 contiguous amino acids of the wild-type sequence). In one embodiment, a homologue of the present invention includes proteins having amino acid sequences comprising at least about
25 30, or at least about 40, or at least about 45, or at least about 50, or at least about 55, or at least about 60, or at least about 65, or at least about 70, or at least about 75, or at least about 80, or at least about 85, or at least about 90, contiguous amino acid residues of the wild-type sequence, and so on, in whole integers up to just less than the full length of the wild-type protein. According to the present invention, the term "contiguous" or "consecutive", with
30 regard to nucleic acid or amino acid sequences described herein, means to be connected in an unbroken sequence. For example, for a first sequence to comprise 30 contiguous (or

consecutive) amino acids of a second sequence, means that the first sequence includes an unbroken sequence of 30 amino acid residues that is 100% identical to an unbroken sequence of 30 amino acid residues in the second sequence. Similarly, for a first sequence to have "100% identity" with a second sequence means that the first sequence exactly matches the second sequence with no gaps between nucleotides or amino acids.

Further, any of the amino acid sequences described herein can be produced with from at least one, and up to about 20, additional heterologous amino acids flanking each of the C- and/or N-terminal ends of the specified amino acid sequence. The resulting protein or polypeptide can be referred to as "consisting essentially of" the specified amino acid sequence. According to the present invention, the heterologous amino acids are a sequence of amino acids that are not naturally found (i.e., not found in nature, *in vivo*) flanking the specified amino acid sequence, or that are not related to the function of the specified amino acid sequence, or that would not be encoded by the nucleotides that flank the naturally occurring nucleic acid sequence encoding the specified amino acid sequence as it occurs in the gene, if such nucleotides in the naturally occurring sequence were translated using standard codon usage for the organism from which the given amino acid sequence is derived. Similarly, the phrase "consisting essentially of", when used with reference to a nucleic acid sequence herein, refers to a nucleic acid sequence encoding a specified amino acid sequence that can be flanked by from at least one, and up to as many as about 60, additional heterologous nucleotides at each of the 5' and/or the 3' end of the nucleic acid sequence encoding the specified amino acid sequence. The heterologous nucleotides are not naturally found (i.e., not found in nature, *in vivo*) flanking the nucleic acid sequence encoding the specified amino acid sequence as it occurs in the natural gene or do not encode a protein that imparts any additional function to the protein or changes the function of the protein having the specified amino acid sequence.

With regard to specific embodiments of the invention, one aspect of the invention relates to a TALL-1 antagonist protein. A TALL-1 antagonist protein is a TALL-1 homologue (i.e., mutant) that antagonizes the biological activity of a wild-type, or naturally occurring TALL-1 protein, as discussed above. A TALL-1 antagonist is generally a TALL-1 homologue which comprises at least one amino acid modification as compared to a naturally occurring TALL-1 or is a portion of TALL-1 that contains the modification. The

modifications to the amino acid sequence of the mutant TALL-1 can include any of the modifications to any amino acid position corresponding to any of the target residues identified herein based on the determination of the structure of TALL-1. In one embodiment of the invention, an antagonist TALL-1 protein is disclosed that has an amino acid sequence comprising at least one modification as compared to a naturally occurring TALL-1, wherein the modification is in a region selected from: (1) the "flap" region of TALL-1 (discussed in detail below); (2) a region other than the "flap" that participates in trimer-trimer associations or "clustering" of TALL-1 trimers; (3) a region that is involved in formation of TALL-1 trimers; and/or (4) a region of TALL-1 that is associated with binding to a TALL-1 receptor (e.g., BCMA, BAFF-R or TACI). Specific regions of TALL-1 and amino acid residues that are associated with each of these functions are described in detail in the Examples section.

In a first embodiment, the TALL-1 antagonist protein comprises an amino acid sequence that differs from the amino acid sequence of the wild-type protein (e.g., SEQ ID NO:2) by a modification in at least one region of the protein or in at least one amino acid residue of the protein that was determined by the present inventors, based on the structural analysis of TALL-1 disclosed herein, to play a role in the biological activity of TALL-1, other than the binding of TALL-1 to its receptor. Resulting TALL-1 antagonist proteins will have reduced biological activity as compared to a wild-type protein and can serve as competitive inhibitors of TALL-1, for example. Preferably, these TALL-1 antagonists will retain the ability to bind to a TALL-1 receptor and in a most preferred embodiment, these TALL-1 antagonists will have modifications that result in *increased* binding affinity for a TALL-1 receptor as compared to the binding affinity between wild-type TALL-1 and the receptor.

In one aspect, the above-described TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by at least one modification in the region connecting β strands D and E that reduces the biological activity of the TALL-1 antagonist as compared to wild-type TALL-1. As discussed in detail above and in the Examples, the present inventors have discovered that TALL-1 has a unique structure denoted the "flap" which mediates the viral-like clustering of TALL-1 trimers and which the inventors have shown is necessary for the biological activity of TALL-1. The region connecting β strands D and E of TALL-1 (see examples and figures) forms this flap, which includes amino acid residues 217-224 of SEQ ID NO:2. The present inventors have also produced a mutant

TALL-1 in which these residues were deleted and replaced with two glycine residues (see Example 3). This mutant, while still able to bind to the TALL-1 receptors, was not able to activate the receptor or stimulate B cell proliferation (i.e., overall, it did not have TALL-1 biological activity). Moreover, the present inventors have shown that this deletion abolished the cluster forming ability of TALL-1 homologue and had a negative effect on the formation of TALL-1 trimers (see Examples). In one embodiment, this TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from: Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224. In another aspect, this TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least two, three, four, five, six, seven, or all eight of the above-identified amino acid residues. It is noted that throughout the application, amino acid residues are denoted using the art-recognized 3 letter code.

According to the invention, the residues can be deleted, derivatized to reduce the ability of the residues to interact with residues on other TALL-1 monomers as described herein for the flap region residues, or substituted with non-natural amino acid residues that reduce the ability of the homologue to interact with residues on other TALL-1 monomers as described herein for the flap region residues. As used throughout this disclosure, reference to "non-natural" residues describes amino acid residues that are used to modify a protein (e.g., by insertion), but which do not naturally occur at the insertion position in the wild-type protein. In addition, one or more non-natural amino acid residues can be inserted to replace one or more (not necessarily an equivalent number) of deleted residues in the homologue sequence. For example, as exemplified herein, two glycine residues were inserted in place of a deletion of the entire 8 amino acid flap sequence to produce a TALL-1 antagonist with reduced biological activity.

Modifications in this region can produce a TALL-1 antagonist protein that has a reduced ability to form a trimer with other TALL-1 monomers. "Other TALL-1 monomers", as used herein, can refer to any TALL-1 monomer other than the monomer that is the reference homologue, including wild-type TALL-1 monomers and/or other TALL-1 homologue monomers (having the same or different modifications as the reference homologue). Therefore, a TALL-1 antagonist monomer can associate (e.g., to form a trimer,

or when already in a trimer, to associate with another trimer), or attempt to associate, with wild-type monomers, with other TALL-1 antagonist monomers, or with mixtures thereof (e.g., to form heterotrimers), *in vitro* or *in vivo*. Modifications in this region can also produce a TALL-1 antagonist that has a reduced or abolished ability, when in a trimer with two other
5 TALL-1 monomers, to interact with other TALL-1 trimers, such as to form the viral-like cluster of trimers described herein. According to the present invention, to reduce binding between a TALL-1 homologue and another protein (e.g., monomers, trimers, receptors) refers to any detectable decrease in the binding affinity as compared to the binding affinity between the wild-type TALL-1 monomer and the same other protein. To abolish binding between a
10 TALL-1 homologue and another protein refers to a substantial abolition (reduction, elimination, prevention) of binding affinity as compared to the binding affinity between the wild-type TALL-1 monomer and the same other protein, wherein the binding affinity is reduced by at least about 50%, and more preferably by at least about 60%, and more preferably by at least about 70%, and more preferably by at least about 80%, and more
15 preferably at least about 90%, and more preferably at least about 95%, and more preferably at least about 96%, and more preferably at least about 97%, and more preferably at least about 98%, and more preferably at least about 99%, and most preferably, wherein binding of the mutated TALL-1 monomer to another protein is undetectable using standard binding assays. Similarly, to increase binding between a TALL-1 antagonist and another protein
20 (e.g., a receptor; see discussion below) refers to any detectable increase in the binding affinity as compared to the binding affinity between the wild-type TALL-1 monomer and the same other protein. Assays for detecting and measuring binding, including binding affinity, between two proteins are well known in the art and have been discussed previously herein.

As discussed above, it is preferred that the above-described TALL-1 antagonist
25 protein retain the ability to bind to a TALL-1 receptor, including, but not limited to, BCMA, BAFF-R and TACI. In this way, the TALL-1 antagonist can competitively inhibit wild-type TALL-1 by binding to, but not activating, TALL-1 receptors. In one aspect, the TALL-1 antagonist protein retain some or all of the amino acid residues that participate in binding to a TALL-1 receptor, including, but not limited to, Tyr163, Tyr206, Leu211, Arg231, Ile233,
30 Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222 of SEQ ID NO:2. The residues are spatially arranged in the amino acid sequence of the TALL-1

homologue in a manner that is similar enough to the spatial arrangement in the wild-type TALL-1, so that binding to a TALL-1 receptor is retained. At a minimum, this includes the amino acid residues that occur in positions 163-275 in the wild-type protein. However, it is to be understood that the positions of these residues within the mutated TALL-1 protein can vary somewhat from the corresponding positions in the wild-type protein, as long as the mutated TALL-1 protein maintains the ability to bind to a TALL-1 receptor. For example, one might be able to construct a TALL-1 antagonist protein where the Tyr that occurs at position 163 of the wild-type protein appears at position 162 or 164 in the mutant. Moreover, to the extent that intervening residues are required to maintain the approximate distance between the critical receptor binding residues, one can construct the mutated TALL-1 protein accordingly, such as by retaining additional wild-type sequence or by using conservative amino acid substitutions in the intervening residues that maintain the three-dimensional structure of the receptor binding site as disclosed herein. In a particularly preferred aspect, the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2 by at least one additional modification (i.e., in addition to the modification to decrease biological activity) that *increases* the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said TALL-1 receptor. Such modifications are preferably made at one or more of the following positions with respect to SEQ ID NO:2: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

In yet another embodiment, a TALL-1 antagonist protein that has reduced biological activity as compared to a wild-type TALL-1 protein includes a protein that comprises an amino acid sequence that differs from SEQ ID NO:2 by at least one modification that reduces interaction between a first trimer and a second trimer. As discussed above, TALL-1 monomers form trimers and the present inventors have shown that these trimers interact to form viral-like clusters that are believed to represent the biologically active form of TALL-1

in vivo. Reduction of the ability of the TALL-1 monomers to participate in trimer-trimer interactions will therefore reduce the biological activity of the protein. In addition to the "flap" region discussed above, the present inventors have identified many other residues in TALL-1 that are involved in the trimer-trimer associations (see Examples). An antagonist TALL-1 protein according to this embodiment may form trimers with any other TALL-1 monomers, including wild-type TALL-1 monomers and other TALL-1 homologue monomers (the same or different than the reference monomer), and the trimers can be homotrimers (formed of monomers of all the same type) or heterotrimers (formed of monomers of different types, such as two wild-type monomers with one antagonist monomer). In one aspect, this antagonist may also have a reduced ability to form trimers with other TALL-1 monomers (wild-type or homologues).

In one aspect of this embodiment of the invention, the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue located in a region of TALL-1 selected from the group consisting of β strand C, β strand F, and the region connecting β strand D to β strand E. In another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from: Ile150, Leu169, Phe172, Tyr192, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Ile250, Lys252, and Glu254. In further embodiments, any number of residues greater than one and up to all of these residues can be modified. In one aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from: Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224. In another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from the group consisting of: Tyr192, Lys252, Glu254, His218, Lys216, Glu223, Leu224, Val227, Leu229, Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172. In another embodiment, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from the group consisting of: Tyr192, Lys252, Glu254, and His218. In yet another embodiment, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from the group consisting of: Lys216,

Glu223, Leu224, Val227, and Leu229. In yet another embodiment, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification in at least one amino acid residue selected from the group consisting of: Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172.

5 As described above for modifications to the flap region, the modified amino acid residues can be deleted, derivatized to reduce the ability of the residues to interact with residues on other TALL-1 monomers as described herein for the trimer-trimer associations, or substituted with non-natural amino acid residues that reduce the ability of the homologue to interact with residues on other TALL-1 monomers as described herein for the trimer-trimer
10 associations. In addition, one or more non-natural amino acid residues can be inserted to replace one or more (not necessarily an equivalent number) of deleted residues in the homologue sequence. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

15 Preferably, this TALL-1 antagonist protein binds to a TALL-1 receptor, including, but not limited to, BCMA, BAFF-R and TACI. In a more preferred embodiment, the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-
20 type TALL-1 and the TALL-1 receptor. For example, as discussed above, the TALL-1 antagonist can comprise an amino acid sequence that differs from SEQ ID NO:2 by an additional modification in at least one amino acid residue selected from the group consisting of: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, where the additional modification increases the
25 binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

30 In another embodiment, a TALL-1 antagonist protein that has reduced biological activity as compared to a wild-type TALL-1 protein includes a protein that comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification of at least one amino

acid residue selected from the group consisting of: Phe194, Tyr196, Tyr246, Leu282, Gln144 and Leu285. According to the present invention, these residues have been identified by the present inventors as being associated with the ability of TALL-1 monomers to form trimers, which is generally agreed in the art to be necessary for TALL-1 activity. A TALL-1 antagonist that has a reduced, or even abolished ability to form trimers, preferably retains the ability to bind to a TALL-1 receptor. The present inventors have shown that TALL-1 and its receptor bind in *mostly* one to one ratio, although a second monomer from a TALL-1 trimer does contribute to the interaction. Therefore, although binding to a single receptor by a monomer may be less stable than binding of a TALL-1 trimer, it is believed that a TALL-1 antagonist according to this embodiment will be able to compete at least to some extent with wild-type TALL-1 for binding to a receptor and thereby act as a competitive inhibitor. Moreover, introduction into this TALL-1 antagonist of additional modifications that increase the binding of the antagonist to the receptor, as described previously herein, can increase the efficacy of this antagonist as an inhibitor. Therefore, in one aspect, this TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor, such residues including, but not limited to: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, where the additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

In yet another embodiment of the invention, a TALL-1 antagonist protein that has reduced biological activity as compared to a wild-type TALL-1 protein includes a protein that comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification of at least one amino acid residue that reduces the biological activity of the antagonist protein as compared to a wild-type TALL-1, wherein the amino acid residue is selected from: Gln144,

Ile150, Leu169, Phe172, Tyr192, Phe194, Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285. This antagonist further differs from SEQ ID NO:2 by a modification of at least one amino acid residue that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and the TALL-1 receptor, wherein the amino acid residue is selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

Yet another embodiment of the present invention relates to a TALL-1 antagonist protein, that has a reduced ability to bind to a receptor for TALL-1, including but not limited to, BCMA, BAFF-R or TACI. In this embodiment, the TALL-1 antagonist may otherwise have substantially normal biological activity as compared to the wild-type TALL-1 protein, or may have reduced biological activity, using the modifications described above, although preferably, this TALL-1 antagonist retains the ability to form trimers and participate in trimer-trimer interactions. This TALL-1 antagonist is useful as an agent the binds to wild-type TALL-1 monomers and thus forms part of trimers and clusters that have a reduced ability to bind to TALL-1 receptors, by virtue of the modifications to the receptor-binding portion of the antagonist. This TALL-1 antagonist is likely to be most efficient if delivered to a cell such that the antagonist is expressed by a cell that also expresses wild-type TALL-1 monomers, such that the antagonist can readily form the "defective" trimers as the proteins are expressed by a cell. In this embodiment, a recombinant nucleic acid molecule encoding the antagonist is delivered to the cell so that the cell expresses the antagonist and so that the antagonist can assemble trimers with wild-type TALL-1 monomers. Alternatively, the TALL-1 antagonist can be delivered to a cell or site under conditions whereby TALL-1 trimers may not have formed the viral-like clusters described herein. Without being bound by theory, the present inventors believe that trimers of TALL-1, prior to assembly into clusters, are in equilibrium, so that a TALL-1 antagonist near such trimers could reassemble

with the wild-type monomers during this equilibrium. Once the viral-like assembly forms, which can be driven by concentration of trimers at a site, for example, the clustered assembly is not believed to be in equilibrium but rather quite stable. Therefore, if trimers can be formed that incorporate the TALL-1 antagonist prior to cluster formation, the result would be stable clusters with overall reduced receptor binding capability, and thus reduced TALL-1 activity via reduced TALL-1 receptor activation.

In one aspect of this embodiment, the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification to at least one amino acid residue selected from: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222, wherein the TALL-1 antagonist protein has reduced binding to a receptor for TALL-1 as compared to wild-type TALL-1. In another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification to at least one amino acid residue selected from: Tyr163, Leu211, Ile233, Pro264, and Leu200. In another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification to at least one amino acid residue selected from the group consisting of: Tyr206 and Leu240. In yet another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification to at least one amino acid residue selected from the group consisting of: Arg265, Glu266 and Glu238. In yet another aspect, the antagonist comprises an amino acid sequence that differs from SEQ ID NO:2 by a modification to at least one amino acid residue selected from the group consisting of: Asp222, Asp 273 and Asp275. In one embodiment, the TALL-1 antagonist has reduced ability to bind to at least two of BCMA, BAFF-R and TACI, and in another embodiment, has a reduced ability to bind to each of BCMA, BAFF-R and TACI. Given the knowledge provided herein of the common and different residues of TALL-1 that are used in receptor binding to at least BCMA and BAFF-R, design and selection of such an antagonist is predicted. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

It is to be expressly understood that any and all of the above-identified amino acid regions and residues that are important for TALL-1 biological activity and receptor binding can also be modified to produce TALL-1 agonists. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues. A TALL-1 agonist is a TALL-1 homologue (i.e., mutant) that is an agonist of the biological activity of a wild-type, or naturally occurring TALL-1 protein, as discussed above. A TALL-1 agonist is generally a TALL-1 homologue which comprises at least one amino acid modification as compared to a naturally occurring TALL-1 or is a portion of TALL-1 that contains the modification, wherein the result is a homologue with TALL-1 activity or increased activity. The modifications to the amino acid sequence of the mutant TALL-1 can include any of the modifications to any amino acid position corresponding to any of the target residues identified herein based on the determination of the structure of TALL-1. In one embodiment of the invention, an agonist TALL-1 protein is disclosed that has an amino acid sequence comprising at least one modification as compared to a naturally occurring TALL-1, wherein the modification is in a region selected from: (1) the "flap" region of TALL-1 (discussed in detail below); (2) a region other than the "flap" that participates in trimer-trimer associations or "clustering" of TALL-1 trimers; (3) a region that is involved in formation of TALL-1 trimers; and/or (4) a region of TALL-1 that is associated with binding to a TALL-1 receptor (e.g., BCMA, BAFF-R or TACI). In contrast to TALL-1 antagonists, TALL-1 agonists with modifications in these regions have the same or increased ability to form trimers or to participate in trimer-trimer interactions, and/or have the same or increased ability to activate TALL-1 receptors, as compared to a wild-type TALL-1 protein. In addition, TALL-1 agonists can bind to TALL-1 receptors and can have increased binding to TALL-1 receptors as compared to the wild-type protein. Specific regions of TALL-1 and amino acid residues that are associated with each of these functions are described in detail in the Examples section.

Another embodiment of the present invention relates to an APRIL agonist protein. As discussed above, the present inventors, without being bound by theory, believe that

APRIL may be serving as a decoy ligand, reducing the opportunity for sTALL-1 to bind to the same receptor. This role is similar to the decoy death receptors, which are essential for cells to survive. Therefore, agonists of APRIL, as well as wild-type APRIL itself, can effectively serve as TALL-1 antagonists according to the present invention. Therefore, the invention contemplates the production of homologues of APRIL which retain the biological activity of APRIL and which preferably bind to APRIL receptors (two of which are shared by TALL-1 - BCMA and TACI). In a more preferred embodiment, the APRIL agonist has an increased ability to bind to an APRIL or TALL-1 receptor, and in another preferred embodiment, the receptor binding site of APRIL is modified so that APRIL binds to TALL-1 receptor such as BAFF-R (see Example 6). In one embodiment, the APRIL agonist protein comprises an amino acid sequence that differs from SEQ ID NO:4 by at least one modification that increases the binding affinity between the APRIL agonist protein and an APRIL receptor, as compared to the binding affinity between wild-type APRIL and the APRIL receptor. For example, in one aspect, such an agonist comprises an amino acid sequence that differs from SEQ ID NO:4 by a modification in at least one amino acid residue selected from: Val133, Thr177, Val181, Ile197, Pro230, Leu58, Tyr96, Phe176, Arg206, and Arg265, where the modification increases the binding affinity between the APRIL agonist protein and an APRIL receptor, as compared to the binding affinity between wild-type APRIL and the APRIL receptor. The modifications can include deletions, derivatizations, and/or substitutions of amino acids and/or insertion of non-natural amino acids effective to achieve the desired result, as discussed above. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues.

In another embodiment, APRIL antagonists are contemplated. APRIL antagonists can have any one or more of the modifications in preferred amino acid residues, or in the regions surrounding these residues as described above for APRIL agonists, except the selected result is a modification that provides a homologue with biological activity that is antagonistic to the biological activity of wild-type APRIL.

Another embodiment of the present invention relates to antagonists of TALL-1 receptors, including antagonists of BCMA and/or BAFF-R, as well as antagonists of TACI. A TALL-1 receptor antagonist is a TALL-1 receptor homologue (i.e., mutant) that

antagonizes the biological activity of a wild-type, or naturally occurring TALL-1 receptor, as discussed above. A TALL-1 receptor antagonist is generally a TALL-1 receptor homologue which comprises at least one amino acid modification as compared to a naturally occurring TALL-1 receptor or is a portion of a TALL-1 receptor that contains the modification. The modifications to the amino acid sequence of the mutant TALL-1 receptor can include any of the modifications to any amino acid position corresponding to any of the target residues identified herein based on the determination of the structure of the extracellular domains of the TALL-1 receptors BCMA and BAFF-R and can extend to TACI, given the amino acid similarity between BCMA and TACI and the prediction of similar structures. In one embodiment of the invention, an antagonist TALL-1 receptor protein is disclosed that has an amino acid sequence comprising at least one modification as compared to a naturally occurring TALL-1 receptor, wherein the modification is in a region that interacts with the natural ligand for the receptor (e.g., TALL-1 or APRIL). Specific regions of TALL-1 and amino acid residues that are associated with ligand binding are described in detail in the Examples section. In a preferred embodiment, the TALL-1 receptor antagonist is a soluble TALL-1 receptor, with the modifications to the amino acid sequence of the TALL-1 receptor, described herein. Soluble TALL-1 receptors, such as soluble BCMA, are described in detail in U.S. Patent Application No. 09/565,423 to Shu, incorporated herein by reference in its entirety.

In one aspect of this embodiment of the invention, the TALL-1 receptor antagonist is a BCMA antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:6 by a modification in at least one amino acid residue selected from: Tyr13, Asp15, Leu17, Leu18, His19, Ile22, Leu26, Arg27, and Pro34, wherein the BCMA antagonist has an increased binding affinity for TALL-1 as compared to wild-type BCMA. In one aspect, the amino acid residue to be modified is selected from Leu17 and Leu18. In another aspect, the amino acid residue to be modified is selected from Ile22 and Leu26. In another aspect, the amino acid residue to be modified is selected from Asp15, Arg27 and Tyr13. In yet another aspect, the amino acid residue to be modified is His19. In yet another aspect, the amino acid residue to be modified is selected Tyr13, Leu17, Leu18 and Ile22. In a preferred embodiment, any one or more of Tyr13, Leu17, Leu18 and Ile22 is substituted with any one of the amino acid residues selected from: Ile, Met, Phe or Tyr.

In each of these embodiments, the receptor antagonist preferably has an increased affinity for a ligand of the receptor (e.g., TALL-1 or APRIL), so that the receptor can serve as a competitive inhibitor of the natural receptor (wild-type). In addition, to serve as an antagonist of the wild-type receptor, the antagonist should have reduced ability to induce a signal that is associated with activation of the wild-type receptor or preferably, the receptor antagonist should not be able to transduce a signal, such as in a soluble receptor.

In another embodiment, the TALL-1 receptor antagonist is a BCMA antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:6 by a modification in at least one amino acid residue within 2-5 amino acid residues to either side of any of the above-identified amino acid residues, including the above-identified amino acid residues. The basic tertiary structure of the BCMA receptor (at least the ligand binding region) should be maintained, which can be readily accomplished given the detailed disclosure of the tertiary structure of eBCMA provided herein.

The modified amino acid residues can be deleted, derivatized to increase the ability of the residues to interact with residues on the receptor ligand (e.g., TALL-1 or APRIL), or substituted with non-natural amino acid residues that increase the ability of the homologue to interact with residues on the receptor ligand. In addition, one or more non-natural amino acid residues can be inserted to replace one or more (not necessarily an equivalent number) of deleted residues in the homologue sequence. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues. Binding affinity for a receptor and ligand can be readily measured as described previously herein.

In another aspect, BCMA homologues are provided with altered binding to APRIL (agonists or antagonists). For example, given the information provided herein (see Example 6), one can produce BCMA homologues with a modification at His19 which has modified binding to APRIL.

In another aspect of this embodiment of the invention, the TALL-1 receptor antagonist is a BAFF-R antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:8 by a modification in at least one amino acid residue selected from: Asp26, Leu28, Val29, Arg30, Val33, Leu37, Leu38, and Arg42, and Pro45, where the BAFF-R antagonist has an increased binding affinity for TALL-1 as compared to

wild-type BAFF-R. In one aspect, the amino acid residue to be modified is selected from Leu28 and Val29. In another aspect, the amino acid residue to be modified is selected from Val33, Leu37, Leu38 and Pro45. In another aspect, the amino acid residue to be modified is selected from Asp26 and Arg 42. In yet another aspect, the amino acid residue to be modified is Arg30. In yet another aspect, the amino acid residue to be modified is selected from Leu28, Val29 and Val33. In another aspect, any one or more of amino acid residues selected from Leu28, Val29 and Val33 is substituted with any one of the following amino acid residues: Ile, Met, Phe or Tyr. In each of these embodiments, the receptor antagonist preferably has an increased affinity for a ligand of the receptor (e.g., TALL-1), so that the receptor can serve as a competitive inhibitor of the natural receptor (wild-type). In addition, to serve as an antagonist of the wild-type receptor, the antagonist should have reduced ability to induce a signal that is associated with activation of the wild-type receptor or preferably, the receptor antagonist should not be able to transduce a signal, such as in a soluble receptor.

In another embodiment, the TALL-1 receptor antagonist is a BAFF-R antagonist, wherein the receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:8 by a modification in at least one amino acid residue within 2-5 amino acid residues to either side of any of the above-identified amino acid residues, including the above-identified amino acid residues. The basic tertiary structure of the BAFF-R (at least the ligand binding region) should be maintained, which can be readily accomplished given the detailed disclosure of the tertiary structure of eBAFF-R provided herein.

In yet another embodiment, a TALL-1 receptor antagonist or agonist (depending on the referenced function of the homologue) of BAFF-R is produced which binds to APRIL. Using the guidance provided in Example 6, for example, one can modify one or more residues in BAFF-R to provide a homologue of BAFF-R that can bind to APRIL. Such residues include, but are not limited to, modification of residues 1-11 or 1-12 of SEQ ID NO:8; modification of Arg30; modification of His31; modification of Val29; and/or modification of Val33. Two mutants of BAFF-R with the ability to bind to APRIL at pH7.5 are described in Example 6 (i.e., (1) deletion of residues 1-11 and Arg30His and His31Arg; (2) deletion of residues 1-11 and Val29Leu and Val33Ile). Other modifications producing similar results will be apparent to those of skill in the art given the structural information for APRIL and BAFF-R provided herein.

The modified amino acid residues can be deleted, derivatized to increase the ability of the residues to interact with residues on the receptor ligand (e.g., TALL-1), or substituted with non-natural amino acid residues that increase the ability of the homologue to interact with residues on the receptor ligand. In addition, one or more non-natural amino acid residues can be inserted to replace one or more (not necessarily an equivalent number) of deleted residues in the homologue sequence. Modifications are made to at least one residue, but can be made to two, three, four, or any additional number of the above-identified residues, up to all of these residues. Binding affinity for a receptor and ligand can be readily measured as described previously herein.

It is to be expressly understood that any and all of the above-identified amino acid regions and residues that are important for TALL-1 receptor-ligand binding can also be modified to produce TALL-1 receptor agonists, if desired. Also, it is to be understood that, using the structural information disclosed herein, one can predict the corresponding residues of TACI that will interact with BCMA or APRIL, and design antagonists of this receptor.

Given the structural information provided herein, one of skill in the art will also be able to design polypeptides of TALL-1 or its receptors which fall within the scope of the embodiments described above and serve as agonists or antagonists of TALL-1 or its receptors. By way of example, yet another embodiment of the present invention relates to an isolated BAFF-R antagonist that consists essentially of the amino acid sequence represented herein by SEQ ID NO:9. Positions 2-26 of SEQ ID NO:9 correspond to amino acid positions 15-29 of SEQ ID NO:8, with the following substitutions. At position 2 of SEQ ID NO:9, a Ser is substituted for the Ala that occurs in BAFF-R (position 15 of SEQ ID NO:8); at position 16 of SEQ ID NO:9, a Leu is substituted for the Val that occurs in BAFF-R (position 29 of SEQ ID NO:8); and at position 19 of SEQ ID NO:9, an Ile is substituted for the Val that occurs in BAFF-R (position 33 of SEQ ID NO:8). Without being bound by theory, the present inventors believe that extension of amino acid side chains in the ligand binding region of the receptor will increase the affinity of the receptor homologue for the ligand.

Another embodiment of the invention relates to a regulatory peptide having the amino acid sequence of SEQ ID NO:10, which is a peptide from the flap region of TALL-1 disclosed herein.

Another embodiment of the invention relates to a regulatory peptide or a homologue thereof having the amino acid sequence of any one of SEQ ID NO:11-SEQ ID NO:16. These sequences are illustrated in Fig. 8B and show the strong pattern of similarity between BCMA (SEQ ID NO:11), BAFF-R (SEQ ID NO:12), TACI1 (SEQ ID NO:13), TACI2 (SEQ ID NO:14), Fn14 (SEQ ID NO:15), and TNF-R1 (SEQ ID NO:16). Shown in Fig. 8B are residues (colored red) that are believed to be conserved disulfide bridges or pseudo disulfide bridges, which builds up module A1, D2, and D0. Residues colored yellow are not defined. Residues colored blue are for the C2 module. Residues colored green are putative residues involved in ligand recognition. Any of these residues, and particularly those that are at or near positions of defined proposed function (e.g., the green residues that are proposed to be involved in ligand recognition), can be targeted for modification in any of these proteins to produce novel homologues of the specified protein with altered biological activity and ligand binding, as described herein in detail for BCMA and BAFF-R.

Another embodiment of the invention relates to a homologue of any of TACI1, TACI2, Fn14, and TNF-R1, wherein the amino acid sequence of the homologue differs from the amino acid sequence of the wild type modifications (the wild-type protein comprising the amino acid sequence shown in SEQ ID NOs:13-16, respectively) by at least one modification to at least one amino acid residue that modifies the biological activity of the protein. In a preferred embodiment, modifications are made to any of the amino acid positions that are associated with the binding of the receptor to its ligand, as identified herein (see Fig. 8). For example, a TACI receptor can be modified at amino acid positions 39, 41, 43, 52 or 53 of the wild-type protein (TACI1; Fig. 8; corresponding to positions 6, 8, 10, 19 and 20 of SEQ ID NO:13, respectively) or at positions 78, 80, 82, 91 or 92 of the wild-type protein (TACI2; Fig. 8; corresponding to positions 8, 10, 12, 21 and 22 of SEQ ID NO:14, respectively). The complete nucleic acid sequence encoding TACI and the amino acid sequence therefore is provided in GenBank Accession No. AF023614, incorporated herein by reference in its entirety. The complete nucleic acid sequence encoding Fn14 and the amino acid sequence therefore is provided in GenBank Accession No. NM_016639, incorporated herein by reference in its entirety. The complete nucleic acid sequence encoding TNF-R1 and the amino acid sequence therefore is provided in GenBank Accession No. M75866, incorporated herein by reference in its entirety.

Another embodiment of the present invention relates to a fusion protein comprising any of the heretofore described proteins and homologues attached to one or more fusion segments. Suitable fusion segments for use with the present invention include, but are not limited to, segments that can: enhance a protein's stability; provide other desirable biological activity (e.g., a therapeutic protein/peptide to be delivered to a site); and/or assist with the purification of a protein (e.g., by affinity chromatography). A suitable fusion segment can be a domain of any size that has the desired function (e.g., imparts increased stability, solubility, biological activity; and/or simplifies purification of a protein). Fusion segments can be joined to amino and/or carboxyl termini of the protein of interest (e.g., a TALL-1 homologue) and can be susceptible to cleavage in order to enable straight-forward recovery of protein of interest. Fusion proteins are preferably produced by culturing a recombinant cell transfected with a nucleic acid molecule that encodes a protein including the fusion segment attached to either the carboxyl and/or amino terminal end of the protein of interest.

Another embodiment of the present invention relates to an isolated nucleic acid molecule that encodes any of the proteins and homologues described herein. In accordance with the present invention, an isolated polynucleotide, or an isolated nucleic acid molecule, is a nucleic acid molecule that has been removed from its natural milieu (i.e., that has been subject to human manipulation), its natural milieu being the genome or chromosome in which the nucleic acid molecule is found in nature. As such, "isolated" does not necessarily reflect the extent to which the nucleic acid molecule has been purified, but indicates that the molecule does not include an entire genome or an entire chromosome in which the nucleic acid molecule is found in nature. An isolated nucleic acid molecule can include a gene or a portion of a gene (e.g., the regulatory region or promoter). An isolated nucleic acid molecule that includes a gene is not a fragment of a chromosome that includes such gene, but rather includes the coding region and regulatory regions associated with the gene, but no additional genes naturally found on the same chromosome. An isolated nucleic acid molecule can also include a specified nucleic acid sequence flanked by (i.e., at the 5' and/or the 3' end of the sequence) additional nucleic acids that do not normally flank the specified nucleic acid sequence in nature (i.e., heterologous sequences). Isolated nucleic acid molecule can include DNA, RNA (e.g., mRNA), or derivatives of either DNA or RNA (e.g., cDNA). Although the phrase "nucleic acid molecule" primarily refers to the physical nucleic acid

molecule and the phrase "nucleic acid sequence" primarily refers to the sequence of nucleotides on the nucleic acid molecule, the two phrases can be used interchangeably, especially with respect to a nucleic acid molecule, or a nucleic acid sequence, being capable of encoding a protein. Preferably, an isolated nucleic acid molecule of the present invention
5 is produced using recombinant DNA technology (e.g., polymerase chain reaction (PCR) amplification, cloning) or chemical synthesis. If the polynucleotide is an oligonucleotide, such as a probe or primer, the oligonucleotide preferably ranges from about 5 to about 50 or about 500 nucleotides, more preferably from about 10 to about 40 nucleotides, and most preferably from about 15 to about 40 nucleotides in length.

10 Isolated nucleic acid molecules can include coding regions and/or regulatory regions (e.g. promoters), and can include nucleic acid sequences that have been modified by nucleotide insertions, deletions, substitutions, and/or inversions in a manner to encode the various homologues of TALL-1, APRIL or the receptors described herein. An isolated nucleic acid molecule can include degeneracies. As used herein, nucleotide degeneracy
15 refers to the phenomenon that one amino acid can be encoded by different nucleotide codons. Thus, the nucleic acid sequence of a nucleic acid molecule that encodes a protein of the present invention can vary due to degeneracies.

Preferably, an isolated nucleic acid molecule of the present invention is produced using recombinant DNA technology (e.g., polymerase chain reaction (PCR) amplification,
20 cloning) or chemical synthesis. A nucleic acid molecule homologue can be produced using a number of methods known to those skilled in the art (see, for example, Sambrook et al., *ibid.*). For example, nucleic acid molecules can be modified using a variety of techniques including, but not limited to, classical mutagenesis techniques and recombinant DNA techniques, such as site-directed mutagenesis, chemical treatment of a nucleic acid molecule
25 to induce mutations, restriction enzyme cleavage of a nucleic acid fragment, ligation of nucleic acid fragments, PCR amplification and/or mutagenesis of selected regions of a nucleic acid sequence, synthesis of oligonucleotide mixtures and ligation of mixture groups to "build" a mixture of nucleic acid molecules and combinations thereof. Nucleic acid molecule homologues can be selected from a mixture of modified nucleic acids by screening
30 for the function of the protein encoded by the nucleic acid and/or by hybridization with a wild-type gene.

The minimum size of a nucleic acid molecule of the present invention is a size sufficient to encode a protein having the desired biological activity, or sufficient to form a probe or oligonucleotide primer that is capable of forming a stable hybrid with the complementary sequence of a nucleic acid molecule encoding the desired protein (e.g., under moderate, high or very high stringency conditions, and preferably under very high stringency conditions). As such, the size of a nucleic acid molecule of the present invention can be dependent on nucleic acid composition and percent homology or identity between the nucleic acid molecule and complementary sequence as well as upon hybridization conditions *per se* (e.g., temperature, salt concentration, and formamide concentration). The minimal size of a nucleic acid molecule that is used as an oligonucleotide primer or as a probe is typically at least about 12 to about 15 nucleotides in length if the nucleic acid molecules are GC-rich and at least about 15 to about 18 bases in length if they are AT-rich. There is no limit, other than a practical limit, on the maximal size of a nucleic acid molecule of the present invention, in that the nucleic acid molecule can include any functional portion of a protein-encoding sequence (e.g., a TALL-1 homologue-encoding sequence).

One embodiment of the present invention relates to a recombinant nucleic acid molecule which comprises the isolated nucleic acid molecule described above which is operatively linked to at least one transcription control sequence. More particularly, according to the present invention, a recombinant nucleic acid molecule typically comprises a recombinant vector and the isolated nucleic acid molecule as described herein. According to the present invention, a recombinant vector is an engineered (i.e., artificially produced) nucleic acid molecule that is used as a tool for manipulating a nucleic acid sequence of choice and/or for introducing such a nucleic acid sequence into a host cell. The recombinant vector is therefore suitable for use in cloning, sequencing, and/or otherwise manipulating the nucleic acid sequence of choice, such as by expressing and/or delivering the nucleic acid sequence of choice into a host cell to form a recombinant cell. Such a vector typically contains heterologous nucleic acid sequences, that is, nucleic acid sequences that are not naturally found adjacent to nucleic acid sequence to be cloned or delivered, although the vector can also contain regulatory nucleic acid sequences (e.g., promoters, untranslated regions) which are naturally found adjacent to nucleic acid sequences of the present invention or which are useful for expression of the nucleic acid molecules of the present invention

(discussed in detail below). The vector can be either RNA or DNA, either prokaryotic or eukaryotic, and typically is a plasmid. The vector can be maintained as an extrachromosomal element (e.g., a plasmid) or it can be integrated into the chromosome of a recombinant host cell, although it is preferred if the vector remain separate from the genome for most applications of the invention. The entire vector can remain in place within a host cell, or under certain conditions, the plasmid DNA can be deleted, leaving behind the nucleic acid molecule of the present invention. An integrated nucleic acid molecule can be under chromosomal promoter control, under native or plasmid promoter control, or under a combination of several promoter controls. Single or multiple copies of the nucleic acid molecule can be integrated into the chromosome. A recombinant vector of the present invention can contain at least one selectable marker.

In one embodiment, a recombinant vector used in a recombinant nucleic acid molecule of the present invention is an expression vector. As used herein, the phrase "expression vector" is used to refer to a vector that is suitable for production of an encoded product (e.g., a protein of interest). In this embodiment, a nucleic acid sequence encoding the product to be produced (e.g., a TALL-1 homologue) is inserted into the recombinant vector to produce a recombinant nucleic acid molecule. The nucleic acid sequence encoding the protein to be produced is inserted into the vector in a manner that operatively links the nucleic acid sequence to regulatory sequences in the vector which enable the transcription and translation of the nucleic acid sequence within the recombinant host cell.

In another embodiment of the invention, the recombinant nucleic acid molecule comprises a viral vector. A viral vector includes an isolated nucleic acid molecule of the present invention integrated into a viral genome or portion thereof, in which the nucleic acid molecule is packaged in a viral coat that allows entrance of DNA into a cell. A number of viral vectors can be used, including, but not limited to, those based on alphaviruses, poxviruses, adenoviruses, herpesviruses, lentiviruses, adeno-associated viruses and retroviruses.

According to the present invention, the phrase "operatively linked" refers to linking a nucleic acid molecule to a transcription control sequence in a manner such that the molecule is able to be expressed when transfected (i.e., transformed, transduced, transfected, conjugated or conducted) into a host cell. Transcription control sequences are sequences

which control the initiation, elongation, or termination of transcription. Particularly important transcription control sequences are those which control transcription initiation, such as promoter, enhancer, operator and repressor sequences. Suitable transcription control sequences include any transcription control sequence that can function in a host cell or
5 organism into which the recombinant nucleic acid molecule is to be introduced.

Recombinant nucleic acid molecules of the present invention can also contain additional regulatory sequences, such as translation regulatory sequences, origins of replication, and other regulatory sequences that are compatible with the recombinant cell. In one embodiment, a recombinant molecule of the present invention, including those which
10 are integrated into the host cell chromosome, also contains secretory signals (i.e., signal segment nucleic acid sequences) to enable an expressed protein to be secreted from the cell that produces the protein. Suitable signal segments include a signal segment that is naturally associated with the protein to be expressed or any heterologous signal segment capable of directing the secretion of the protein according to the present invention. In another
15 embodiment, a recombinant molecule of the present invention comprises a leader sequence to enable an expressed protein to be delivered to and inserted into the membrane of a host cell. Suitable leader sequences include a leader sequence that is naturally associated with the protein, or any heterologous leader sequence capable of directing the delivery and insertion of the protein to the membrane of a cell.

20 According to the present invention, the term "transfection" is used to refer to any method by which an exogenous nucleic acid molecule (i.e., a recombinant nucleic acid molecule) can be inserted into a cell. The term "transformation" can be used interchangeably with the term "transfection" when such term is used to refer to the introduction of nucleic acid molecules into microbial cells or plants. In microbial systems, the term "transformation"
25 is used to describe an inherited change due to the acquisition of exogenous nucleic acids by the microorganism and is essentially synonymous with the term "transfection." However, in animal cells, transformation has acquired a second meaning which can refer to changes in the growth properties of cells in culture after they become cancerous, for example. Therefore, to avoid confusion, the term "transfection" is preferably used with regard to the introduction
30 of exogenous nucleic acids into animal cells, and is used herein to generally encompass transfection of animal cells and transformation of plant cells and microbial cells, to the extent

that the terms pertain to the introduction of exogenous nucleic acids into a cell. Therefore, transfection techniques include, but are not limited to, transformation, particle bombardment, electroporation, microinjection, lipofection, adsorption, infection and protoplast fusion.

One or more recombinant molecules of the present invention can be used to produce
5 an encoded product (e.g., a TALL-1 antagonist protein, including fusion proteins) of the present invention. In one embodiment, an encoded product is produced by expressing a nucleic acid molecule as described herein under conditions effective to produce the protein. A preferred method to produce an encoded protein is by transfecting a host cell with one or more recombinant molecules to form a recombinant cell. Suitable host cells to transfect
10 include, but are not limited to, any bacterial, fungal (e.g., yeast), insect, plant or animal cell that can be transfected. Host cells can be either untransfected cells or cells that are already transfected with at least one other recombinant nucleic acid molecule.

In one embodiment, one or more protein(s) expressed by an isolated nucleic acid molecule of the present invention are produced by culturing a cell that expresses the protein
15 (i.e., a recombinant cell or recombinant host cell) under conditions effective to produce the protein. In some instances, the protein may be recovered, and in others, the cell may be harvested in whole (e.g., for *ex vivo* administration), either of which can be used in a composition. A preferred cell to culture is any suitable host cell as described above. Effective culture conditions include, but are not limited to, effective media, bioreactor,
20 temperature, pH and oxygen conditions that permit protein production and/or recombination. An effective medium refers to any medium in which a given host cell is typically cultured. Such medium typically comprises an aqueous medium having assimilable carbon, nitrogen and phosphate sources, and appropriate salts, minerals, metals and other nutrients, such as vitamins. Cells can be cultured in conventional fermentation bioreactors, shake flasks, test
25 tubes, microtiter dishes, and petri plates. Culturing can be carried out at a temperature, pH and oxygen content appropriate for a recombinant cell. Such culturing conditions are within the expertise of one of ordinary skill in the art.

Depending on the vector and host system used for production, resultant proteins of the present invention may either remain within the recombinant cell; be secreted into the
30 culture medium; be secreted into a space between two cellular membranes; or be retained on the outer surface of a cell membrane. The phrase "recovering the protein" refers to collecting

the whole culture medium containing the protein and need not imply additional steps of separation or purification. Proteins produced according to the present invention can be purified using a variety of standard protein purification techniques, such as, but not limited to, affinity chromatography, ion exchange chromatography, filtration, electrophoresis, 5 hydrophobic interaction chromatography, gel filtration chromatography, reverse phase chromatography, concanavalin A chromatography, chromatofocusing and differential solubilization.

Another embodiment of the present invention relates to compositions comprising any of the proteins (including homologues) and/or nucleic acid molecules described herein, or 10 any of the compounds identified by drug design/selection using the structures of the present invention, which are useful for screening or therapeutic purposes. Such a composition of the present invention can include any carrier with which the protein, nucleic acid molecule or compound is associated by virtue of the protein, nucleic acid molecule or compound preparation method, a purification method, or a preparation of the protein, nucleic acid 15 molecule or compound for use in an *in vitro*, *ex vivo*, or *in vivo* method according to the present invention. For example, such a carrier can include any suitable excipient, buffer and/or delivery vehicle, such as a pharmaceutically acceptable carrier (discussed below), which is suitable for combining with the protein, nucleic acid molecule or compound of the present invention so that the protein, nucleic acid molecule or compound can be used *in vitro*, 20 *ex vivo* or *in vivo* according to the present invention.

The composition typically also includes a pharmaceutically acceptable carrier. According to the present invention, a "pharmaceutically acceptable carrier" includes pharmaceutically acceptable excipients and/or pharmaceutically acceptable delivery vehicles, which are suitable for use in administration of the composition to a suitable *in vitro*, *ex vivo* 25 or *in vivo* site. A suitable *in vitro*, *in vivo* or *ex vivo* site is preferably a monocyte or macrophage, when TALL-1 is the target molecule (i.e., the molecule which is to be regulated or otherwise targeted by the composition), and a B lymphocyte, when a TALL-1 receptor is the target. In some embodiments, a suitable site for delivery is a site of interaction between B lymphocytes and monocytes or macrophages. Preferred pharmaceutically acceptable 30 carriers are capable of maintaining a protein, compound, or recombinant nucleic acid molecule of the present invention in a form that, upon arrival of the protein, compound, or

recombinant nucleic acid molecule at the cell target in a culture or in patient, the protein, compound or recombinant nucleic acid molecule is capable of interacting with its target (e.g., a naturally occurring TALL-1 protein, including membrane and/or soluble TALL-1 proteins, or a TALL-1 receptor).

5 Suitable excipients of the present invention include excipients or formulations that transport or help transport, but do not specifically target a composition to a cell (also referred to herein as non-targeting carriers). Examples of pharmaceutically acceptable excipients include, but are not limited to water, phosphate buffered saline, Ringer's solution, dextrose solution, serum-containing solutions, Hank's solution, other aqueous physiologically
10 balanced solutions, oils, esters and glycols. Aqueous carriers can contain suitable auxiliary substances required to approximate the physiological conditions of the recipient, for example, by enhancing chemical stability and isotonicity. Compositions of the present invention can be sterilized by conventional methods and/or lyophilized.

One type of pharmaceutically acceptable carrier includes a controlled release
15 formulation that is capable of slowly releasing a composition of the present invention into a patient or culture. As used herein, a controlled release formulation comprises a compound of the present invention (e.g., a protein (including homologues), an antibody, a nucleic acid molecule, or a mimetic) in a controlled release vehicle. Suitable controlled release vehicles include, but are not limited to, biocompatible polymers, other polymeric matrices, capsules,
20 microcapsules, microparticles, bolus preparations, osmotic pumps, diffusion devices, liposomes, lipospheres, and transdermal delivery systems. Other carriers of the present invention include liquids that, upon administration to a patient, form a solid or a gel *in situ*. Preferred carriers are also biodegradable (i.e., bioerodible). When the compound is a recombinant nucleic acid molecule, suitable delivery vehicles include, but are not limited to
25 liposomes, viral vectors or other delivery vehicles, including ribozymes. Natural lipid-containing delivery vehicles include cells and cellular membranes. Artificial lipid-containing delivery vehicles include liposomes and micelles. A delivery vehicle of the present invention can be modified to target to a particular site in a patient, thereby targeting and making use of a compound of the present invention at that site. Suitable modifications include
30 manipulating the chemical formula of the lipid portion of the delivery vehicle and/or introducing into the vehicle a targeting agent capable of specifically targeting a delivery

vehicle to a preferred site, for example, a preferred cell type. Other suitable delivery vehicles include gold particles, poly-L-lysine/DNA-molecular conjugates, and artificial chromosomes.

A pharmaceutically acceptable carrier which is capable of targeting is herein referred to as a "delivery vehicle." Delivery vehicles of the present invention are capable of delivering a composition of the present invention to a target site in a patient. A "target site" refers to a site in a patient to which one desires to deliver a composition. For example, a target site can be any cell which is targeted by direct injection or delivery using liposomes, viral vectors or other delivery vehicles, including ribozymes. Examples of delivery vehicles include, but are not limited to, artificial and natural lipid-containing delivery vehicles, viral vectors, and ribozymes. Natural lipid-containing delivery vehicles include cells and cellular membranes. Artificial lipid-containing delivery vehicles include liposomes and micelles. A delivery vehicle of the present invention can be modified to target to a particular site in a mammal, thereby targeting and making use of a compound of the present invention at that site. Suitable modifications include manipulating the chemical formula of the lipid portion of the delivery vehicle and/or introducing into the vehicle a compound capable of specifically targeting a delivery vehicle to a preferred site, for example, a preferred cell type. Specifically, targeting refers to causing a delivery vehicle to bind to a particular cell by the interaction of the compound in the vehicle to a molecule on the surface of the cell. Suitable targeting compounds include ligands capable of selectively (i.e., specifically) binding another molecule at a particular site. Examples of such ligands include antibodies, antigens, receptors and receptor ligands. Manipulating the chemical formula of the lipid portion of the delivery vehicle can modulate the extracellular or intracellular targeting of the delivery vehicle. For example, a chemical can be added to the lipid formula of a liposome that alters the charge of the lipid bilayer of the liposome so that the liposome fuses with particular cells having particular charge characteristics.

One preferred delivery vehicle of the present invention is a liposome. A liposome is capable of remaining stable in an animal for a sufficient amount of time to deliver a nucleic acid molecule described in the present invention to a preferred site in the animal. A liposome, according to the present invention, comprises a lipid composition that is capable of delivering a nucleic acid molecule described in the present invention to a particular, or selected, site in a patient. A liposome according to the present invention comprises a lipid

composition that is capable of fusing with the plasma membrane of the targeted cell to deliver a nucleic acid molecule into a cell. Suitable liposomes for use with the present invention include any liposome. Preferred liposomes of the present invention include those liposomes commonly used in, for example, gene delivery methods known to those of skill
5 in the art. More preferred liposomes comprise liposomes having a polycationic lipid composition and/or liposomes having a cholesterol backbone conjugated to polyethylene glycol. Complexing a liposome with a nucleic acid molecule of the present invention can be achieved using methods standard in the art.

Another preferred delivery vehicle comprises a viral vector. A viral vector includes
10 an isolated nucleic acid molecule useful in the present invention, in which the nucleic acid molecules are packaged in a viral coat that allows entrance of DNA into a cell. A number of viral vectors can be used, including, but not limited to, those based on alphaviruses, poxviruses, adenoviruses, herpesviruses, lentiviruses, adeno-associated viruses and retroviruses.

15 The various agonists and antagonists described herein (including the TALL-1, APRIL and TALL-1 receptor agonists and antagonists described above and the products of drug design described below) can be used in various therapeutic methods to regulate the biological activity of TALL-1, APRIL, or a TALL-1 receptor. For example, one embodiment of the present invention relates to a method to inhibit TALL-1 biological activity in a mammal,
20 comprising administering to the mammal the recombinant nucleic acid molecule encoding a TALL-1 antagonist protein as described herein, wherein the protein is expressed by a host cell in the mammal. In one aspect of this embodiment, where the TALL-1 antagonist protein has reduced biological activity as compared to wild-type TALL-1, the antagonist associates with wild-type TALL-1 monomers expressed by the cell to produce TALL-1 trimers
25 containing the protein with reduced TALL-1 biological activity, as compared to a trimer of wild-type TALL-1 monomers. In another aspect, where the TALL-1 antagonist has reduced ability to bind to a TALL-1 receptor, the protein associates with wild-type TALL-1 monomers expressed by the cell to produce TALL-1 trimers containing the protein with reduced ability to bind to a TALL-1 receptor, as compared to a trimer of wild-type TALL-1
30 monomers.

Another embodiment relates to a method to inhibit TALL-1 biological activity in a mammal, comprising administering to the mammal any of the TALL-1 antagonist proteins described herein. Preferably, the protein is a competitive inhibitor of wild-type TALL-1 for binding to a TALL-1 receptor.

5 Another embodiment of the invention relates to a method to inhibit TALL-1 receptor biological activity in a mammal, comprising administering to the mammal a TALL-1 receptor antagonist as described herein. In this embodiment, the antagonist is preferably a competitive inhibitor of a wild-type TALL-1 receptor for binding to TALL-1, such as a modified soluble receptor.

10 Another embodiment of the invention relates to a method to inhibit the biological activity of TALL-1, comprising administering to a cell that expresses TALL-1 a recombinant nucleic acid molecule comprising a nucleic acid sequence encoding APRIL, or a biologically active fragment thereof. In this embodiment, a recombinant nucleic acid molecule encoding: wild-type APRIL (SEQ ID NO:4), a biologically active fragment thereof, or a homologue
15 thereof, such as the APRIL agonist described herein, is administered to a cell and preferably, a cell that expresses TALL-1, or a cell near a site where TALL-1 acts, so that the APRIL protein can effectively act as an inhibitor of TALL-1 by binding to the receptor for TALL-1.

According to the present invention, the therapeutic methods of the present invention are primarily directed to the regulation of the biological activity of a target cell (i.e., a B
20 lymphocyte, a monocyte or a macrophage) in a patient with the added, but not required, goal of providing some therapeutic benefit to a patient. Modulating the phenotype of a target cell in a patient in the absence of obtaining some therapeutic benefit is useful for the purposes of determining factors involved (or not involved) in a disease and preparing a patient to more beneficially receive another therapeutic composition. In a preferred embodiment, however,
25 the methods of the present invention are directed to the modulation of the phenotype of a target cell which is useful in providing some therapeutic benefit to a patient. As such, a therapeutic benefit is not necessarily a cure for a particular disease or condition, but rather, preferably encompasses a result which can include alleviation of the disease or condition, elimination of the disease or condition, reduction of a symptom associated with the disease
30 or condition, prevention or alleviation of a secondary disease or condition resulting from the occurrence of a primary disease or condition, and/or prevention of the disease or condition.

As used herein, the phrase "protected from a disease" refers to reducing the symptoms of the disease; reducing the occurrence of the disease, and/or reducing the severity of the disease. Protecting a patient can refer to the ability of a therapeutic composition of the present invention, when administered to a patient, to prevent a disease from occurring and/or to cure or to treat the disease by alleviating disease symptoms, signs or causes. As such, to protect
5 a patient from a disease includes both preventing disease occurrence (prophylactic treatment) and treating a patient that has a disease or that is experiencing initial symptoms or later stage symptoms of a disease (therapeutic treatment). In particular, protecting a patient from a disease or enhancing another therapy (e.g., vaccination) is accomplished by regulating the
10 interaction between TALL-1 and TALL-1 receptor such that a beneficial effect is obtained. A beneficial effect can easily be assessed by one of ordinary skill in the art and/or by a trained clinician who is treating the patient. The term, "disease" refers to any deviation from the normal health of a mammal and includes a state when disease symptoms are present, as well as conditions in which a deviation (e.g., infection, gene mutation, genetic defect, etc.)
15 has occurred, but symptoms are not yet manifested.

In one embodiment, by performing the method of the present invention, the interaction between TALL-1 and TALL-1 receptor is decreased (e.g., using a TALL-1 or TALL-1 receptor antagonist or an APRIL agonist), such a decrease being sufficient to downregulate B lymphocyte proliferation, activation and/or survival in a patient (or in a
20 culture, if the method is performed *in vitro* or *ex vivo*). In one embodiment, when the target cell is an autoreactive B lymphocyte, typically, the patient has or is at risk of developing an autoimmune disease associated with the autoreactive B lymphocyte. Such autoimmune diseases can be any autoimmune disease, and particularly include, rheumatoid arthritis, systemic lupus erythematosus, insulin dependent diabetes mellitus, multiple sclerosis,
25 myasthenia gravis, Grave's disease, autoimmune hemolytic anemia, autoimmune thrombocytopenia purpura, Goodpasture's syndrome, pemphigus vulgaris, acute rheumatic fever, post-streptococcal glomerulonephritis, or polyarteritis nodosa. The autoreactive B lymphocyte in such a patient, prior to the step of administering the composition of the present invention, generally has normal or enhanced proliferation, activation, and/or survival as
30 compared to a B lymphocyte from a patient that does not have and is not at risk of developing the autoimmune disease.

Inhibition of the interaction between TALL-1 and TALL-1 receptor expressed by an autoreactive B lymphocyte can result in a reduction in the proliferation, activation and/or survival of the B lymphocyte, which can be detected as a change in: B lymphocyte cytokine production, a reduction in NF κ B activation, a reduction in TRAF5, TRAF6, NIK, IKK α and IKK β activation, a reduction in immunoglobulin maturation, a reduction in immunoglobulin production and secretion, a reduction in calcium mobilization, or a reduction in phosphorylation of signal transduction proteins. Preferably, inhibition of the interaction between TALL-1 and TALL-1 receptor in the B lymphocytes of the patient produces a result in the patient which includes, but is not limited to, decreased autoantibody production, decreased autoreactive B cell proliferation, decreased autoreactive B cell survival, and/or reduced destruction of autologous cells or tissues, as compared to any of these measurements prior to the conducting of the method of the present invention, or as compared to a patient with the disease who has not been administered the composition of the present invention.

In one embodiment, by performing the method of the present invention, the interaction between TALL-1 and TALL-1 receptor is increased (e.g., by using TALL-1 agonists), such an increase being sufficient to upregulate B lymphocyte proliferation, activation and/or survival in a patient (or in a culture, if the method is performed *in vitro* or *ex vivo*). In one embodiment, the target cell is a normal B lymphocyte (e.g., in a patient receiving a vaccination), an anergic B lymphocyte, or a B lymphocyte in a patient suffering from a suppressed humoral immune response (e.g., in an immune compromised patient). The B lymphocyte in such a patient, prior to the step of administering the composition of the present invention, generally has normal or reduced proliferation, activation, and/or survival as compared to a B lymphocyte from a normal individual, to a patient who is not immune compromised, or to a patient that does not have and is not at risk of developing the disease.

Increasing the interaction between TALL-1 and TALL-1 receptor expressed by a normal or suppressed B lymphocyte can result in an increase in the proliferation, activation and/or survival of the B lymphocyte, which can be detected as a change in: B lymphocyte cytokine production, an increase in NF κ B activation, an increase in TRAF5, TRAF6, NIK, IKK α and IKK β activation, an increase in immunoglobulin maturation, an increase in immunoglobulin production and secretion, an increase in calcium mobilization, and/or an increase in phosphorylation of intracellular signal transduction proteins. Preferably,

increasing the interaction between TALL-1 and TALL-1 receptor in the B lymphocytes of the patient produces a result in the patient which includes, but is not limited to, increased antibody production, increased B cell proliferation, and increased B cell survival, as compared to any of these measurements prior to the conducting of the method of the present invention, or as compared to a patient with the disease who has not been administered the composition of the present invention.

More specifically, a therapeutic composition as described herein, when administered to a patient by the method of the present invention, preferably produces a result which can include alleviation of the disease (e.g., reduction of at least one symptom or clinical manifestation of the disease), elimination of the disease, reduction in inflammation associated with the disease, increased clearance of infectious organisms associated with the disease, reduction of a tumor or lesion associated with the disease, elimination of a tumor or lesion associated with the disease, prevention or alleviation of a secondary disease resulting from the occurrence of a primary disease, prevention of the disease, and initial control or induction of effector cell immunity and/or humoral immunity (i.e., adaptive immunity) against the disease.

According to the present invention, an effective administration protocol (i.e., administering a therapeutic composition in an effective manner) comprises suitable dose parameters and modes of administration that result in the desired effect in the patient (e.g., regulation of B cell proliferation, activation and/or survival), preferably so that the patient is protected from the disease (e.g., by disease prevention or by alleviating one or more symptoms of ongoing disease). Effective dose parameters can be determined using methods standard in the art for a particular disease. Such methods include, for example, determination of survival rates, side effects (i.e., toxicity) and progression or regression of disease.

In accordance with the present invention, a suitable single dose size is a dose that results in regulation of B lymphocyte proliferation, activation and/or survival in a patient, or in the amelioration of at least one symptom of a condition in the patient, when administered one or more times over a suitable time period. Doses can vary depending upon the disease being treated. One of skill in the art can readily determine appropriate single dose sizes for a given patient based on the size of a patient and the route of administration. One of skill in the art can monitor the effectiveness of the treatment by measuring, for example,

determination of survival rates, side effects (i.e., toxicity), determination of cellular and humoral immune response effects, and/or effects on conditions related to B lymphocyte proliferation, activation and/or survival, and symptoms associated with a specific disease or condition.

5 As discussed above, a therapeutic composition of the present invention is administered to a patient in a manner effective to deliver the composition to a cell, a tissue, and/or systemically to the patient, whereby the desired result (e.g., regulation of B lymphocyte proliferation, activation and/or survival) is achieved as a result of the administration of the composition. Suitable administration protocols include any *in vivo* or
10 *ex vivo* administration protocol. The preferred routes of administration will be apparent to those of skill in the art, depending on the type of condition to be prevented or treated; whether the composition is nucleic acid based, protein based, or cell based; and/or the target cell/tissue. For proteins or nucleic acid molecules, preferred methods of *in vivo* administration include, but are not limited to, intravenous administration, intraperitoneal
15 administration, intramuscular administration, intranodal administration, intracoronary administration, intraarterial administration (e.g., into a carotid artery), subcutaneous administration, transdermal delivery, intratracheal administration, subcutaneous administration, intraarticular administration, intraventricular administration, inhalation (e.g., aerosol), intracranial, intraspinal, intraocular, intranasal, oral, bronchial, rectal, topical,
20 vaginal, urethral, pulmonary administration, impregnation of a catheter, and direct injection into a tissue. Routes useful for deliver to mucosal tissues include, bronchial, intradermal, intramuscular, intranasal, other inhalatory, rectal, subcutaneous, topical, transdermal, vaginal and urethral routes. Combinations of routes of delivery can be used and in some instances, may enhance the therapeutic effects of the composition.

25 *Ex vivo* administration refers to performing part of the regulatory step outside of the patient, such as administering a composition (nucleic acid or protein) of the present invention to a population of cells removed from a patient under conditions such that the composition contacts and/or enters the cell, and returning the cells to the patient. *Ex vivo* methods are particularly suitable when the target cell type can easily be removed from and returned to the
30 patient.

Many of the above-described routes of administration, including intravenous, intraperitoneal, intradermal, and intramuscular administrations can be performed using methods standard in the art. Aerosol (inhalation) delivery can also be performed using methods standard in the art (see, for example, Stribling et al., *Proc. Natl. Acad. Sci. USA* 5 189:11277-11281, 1992, which is incorporated herein by reference in its entirety). Oral delivery can be performed by complexing a therapeutic composition of the present invention to a carrier capable of withstanding degradation by digestive enzymes in the gut of an animal. Examples of such carriers, include plastic capsules or tablets, such as those known in the art.

One method of local administration is by direct injection. Direct injection techniques 10 are particularly useful for administering a composition to a cell or tissue that is accessible by surgery, and particularly, on or near the surface of the body. Administration of a composition locally within the area of a target cell refers to injecting the composition centimeters and preferably, millimeters from the target cell or tissue.

Various methods of administration and delivery vehicles disclosed herein have been 15 shown to be effective for delivery of a nucleic acid molecule to a target cell, whereby the nucleic acid molecule transfected the cell and was expressed. In many studies, successful delivery and expression of a heterologous gene was achieved in preferred cell types and/or using preferred delivery vehicles and routes of administration of the present invention. All of the publications discussed below and elsewhere herein with regard to gene delivery and 20 delivery vehicles are incorporated herein by reference in their entirety.

For example, using liposome delivery, U.S. Patent No. 5,705,151, issued January 6, 1998, to Dow et al. demonstrated the successful *in vivo* intravenous delivery of a nucleic acid molecule encoding a superantigen and a nucleic acid molecule encoding a cytokine in a cationic liposome delivery vehicle, whereby the encoded proteins were expressed in tissues 25 of the animal, and particularly in pulmonary tissues. In addition, Liu et al., *Nature Biotechnology* 15:167, 1997, demonstrated that intravenous delivery of cholesterol-containing cationic liposomes containing genes preferentially targets pulmonary tissues and effectively mediates transfer and expression of the genes *in vivo*. Several publications by Dzau and collaborators demonstrate the successful *in vivo* delivery and expression of a gene 30 into cells of the heart, including cardiac myocytes and fibroblasts and vascular smooth muscle cells using both naked DNA and Hemagglutinating virus of Japan-liposome delivery,

administered by both incubation within the pericardium and infusion into a coronary artery (intracoronary delivery) (See, for example, Aoki et al., 1997, *J. Mol. Cell, Cardiol.* 29:949-959; Kaneda et al., 1997, *Ann N.Y. Acad. Sci.* 811:299-308; and von der Leyen et al., 1995, *Proc Natl Acad Sci USA* 92:1137-1141).

5 Delivery of numerous nucleic acid sequences has been accomplished by administration of viral vectors encoding the nucleic acid sequences. Using such vectors, successful delivery and expression has been achieved using *ex vivo* delivery (See, of many examples, retroviral vector; Blaese et al., 1995, *Science* 270:475-480; Bordignon et al., 1995, *Science* 270:470-475), nasal administration (CFTR-adenovirus-associated vector),
10 intracoronary administration (adenoviral vector and Hemagglutinating virus of Japan, see above), intravenous administration (adeno-associated viral vector; Koeberl et al., 1997, *Proc Natl Acad Sci USA* 94:1426-1431). A publication by Maurice et al. (1999, *J. Clin. Invest.* 104:21-29) demonstrated that an adenoviral vector encoding a β 2-adrenergic receptor, administered by intracoronary delivery, resulted in diffuse multichamber myocardial
15 expression of the gene *in vivo*, and subsequent significant increases in hemodynamic function and other improved physiological parameters. Levine et al. describe *in vitro*, *ex vivo* and *in vivo* delivery and expression of a gene to human adipocytes and rabbit adipocytes using an adenoviral vector and direct injection of the constructs into adipose tissue (Levine et al., 1998, *J. Nutr. Sci. Vitaminol.* 44:569-572).

20 In the area of neuronal gene delivery, multiple successful *in vivo* gene transfers have been reported. Millecamps et al. reported the targeting of adenoviral vectors to neurons using neuron restrictive enhancer elements placed upstream of the promoter for the transgene (phosphoglycerate promoter). Such vectors were administered to mice and rats intramuscularly and intracerebrally, respectively, resulting in successful neuronal-specific
25 transfection and expression of the transgene *in vivo* (Millecamps et al., 1999, *Nat. Biotechnol.* 17:865-869). As discussed above, Bennett et al. reported the use of adeno-associated viral vector to deliver and express a gene by subretinal injection in the neural retina *in vivo* for greater than 1 year (Bennett, 1999, *ibid.*).

Gene delivery to synovial lining cells and articular joints has had similar successes.
30 Oligino and colleagues report the use of a herpes simplex viral vector which is deficient for the immediate early genes, ICP4, 22 and 27, to deliver and express two different receptors

in synovial lining cells *in vivo* (Oligino et al., 1999, *Gene Ther.* 6:1713-1720). The herpes vectors were administered by intraarticular injection. Kuboki et al. used adenoviral vector-mediated gene transfer and intraarticular injection to successfully and specifically express a gene in the temporomandibular joints of guinea pigs *in vivo* (Kuboki et al., 1999, *Arch. Oral Biol.* 44:701-709). Apparailly and colleagues systemically administered adenoviral vectors encoding IL-10 to mice and demonstrated successful expression of the gene product and profound therapeutic effects in the treatment of experimentally induced arthritis (Apparailly et al., 1998, *J. Immunol.* 160:5213-5220). In another study, murine leukemia virus-based retroviral vector was used to deliver (by intraarticular injection) and express a human growth hormone gene both *ex vivo* and *in vivo* (Ghivizzani et al., 1997, *Gene Ther.* 4:977-982). This study showed that expression by *in vivo* gene transfer was at least equivalent to that of the *ex vivo* gene transfer. As discussed above, Sawchuk et al. has reported successful *in vivo* adenoviral vector delivery of a gene by intraarticular injection, and prolonged expression of the gene in the synovium by pretreatment of the joint with anti-T cell receptor monoclonal antibody (Sawchuk et al., 1996, *ibid.* Finally, it is noted that *ex vivo* gene transfer of human interleukin-1 receptor antagonist using a retrovirus has produced high level intraarticular expression and therapeutic efficacy in treatment of arthritis, and is now entering FDA approved human gene therapy trials (Evans and Robbins, 1996, *Curr. Opin. Rheumatol.* 8:230-234). Therefore, the state of the art in gene therapy has led the FDA to consider human gene therapy an appropriate strategy for the treatment of at least arthritis. Taken together, all of the above studies in gene therapy indicate that delivery and expression of a recombinant nucleic acid molecule according to the present invention is feasible.

Another method of delivery of recombinant molecules is in a non-targeting carrier (e.g., as "naked" DNA molecules, such as is taught, for example in Wolff et al., 1990, *Science* 247, 1465-1468). Such recombinant nucleic acid molecules are typically injected by direct or intramuscular administration. Recombinant nucleic acid molecules to be administered by naked DNA administration include an isolated nucleic acid molecule of the present invention, and preferably includes a recombinant molecule of the present invention that preferably is replication, or otherwise amplification, competent. A naked nucleic acid reagent of the present invention can comprise one or more nucleic acid molecules of the present invention including a dicistronic recombinant molecule. Naked nucleic acid delivery

can include intramuscular, subcutaneous, intradermal, transdermal, intranasal and oral routes of administration, with direct injection into the target tissue being most preferred. A preferred single dose of a naked nucleic acid vaccine ranges from about 1 nanogram (ng) to about 100 µg, depending on the route of administration and/or method of delivery, as can be
5 determined by those skilled in the art. Suitable delivery methods include, for example, by injection, as drops, aerosolized and/or topically. In one embodiment, pure DNA constructs cover the surface of gold particles (1 to 3 µm in diameter) and are propelled into skin cells or muscle with a "gene gun."

In the method of the present invention, vaccines and therapeutic compositions can be
10 administered to any member of the Vertebrate class, Mammalia, including, without limitation, primates, rodents, livestock and domestic pets. Livestock include mammals to be consumed or that produce useful products (e.g., sheep for wool production). Preferred mammals to protect include humans, dogs, cats, mice, rats, sheep, cattle, horses and pigs, with humans and dogs being particularly preferred, and humans being most preferred.

15 Conditions to treat using methods of the present invention include any condition, disease in which it is useful to modulate the activity of TALL-1 or its receptor(s). Such conditions include, but are not limited to, any condition in which B lymphocyte, monocyte or macrophage activity can be regulated to provide a therapeutic benefit and preferably, includes diseases characterized by hyperproliferation or hypoproliferation of B lymphocytes
20 or hyperactive or hypoactive B cell development, and in one embodiment, diseases characterized by increased numbers of mature B-lymphocytes, splenomegaly, anti-DNA antibodies, proteinuria, or glomerulonephritis. Such conditions include autoimmune disease, such as rheumatoid arthritis, systemic lupus erythematosus, insulin dependent diabetes mellitus, multiple sclerosis, myasthenia gravis, Grave's disease, autoimmune hemolytic
25 anemia, autoimmune thrombocytopenia purpura, Goodpasture's syndrome, pemphigus vulgaris, acute rheumatic fever, post-streptococcal glomerulonephritis, or polyarteritis nodosa. Such diseases also include conditions in which the target cell is a normal B lymphocyte (e.g., in a patient receiving a vaccination), an anergic B lymphocyte, or a B lymphocyte in a patient suffering from a suppressed humoral immune response (e.g., in an
30 immune compromised patient).

Another embodiment of the present invention relates to a method to identify a compound that is a competitive inhibitor of TALL-1 binding to its receptor. The method includes the steps of (a) contacting a TALL-1 receptor or a TALL-1 binding fragment thereof with a homologue of a TALL-1 protein, wherein the homologue comprises an amino acid sequence with a modification in at least one amino acid residue selected from the group consisting of Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, and Glu238; and (b) detecting whether the homologue binds to the TALL-1 receptor or fragment thereof. Homologues that bind to the TALL-1 receptor or fragment thereof potential competitive inhibitors for binding of wild-type TALL-1 to its receptor. The method can further include a step (c) of detecting whether homologues that bind to the TALL-1 receptor or fragment thereof in (b) have a TALL-1 biological activity selected from the group consisting of: an ability to activate signal transduction in the TALL-1 receptor, an ability to form a trimer with two other TALL-1 monomers, an ability to form a trimer with TALL-1 two other TALL-1 monomers that is capable of interacting with other TALL-1 trimers. Homologues that have a decreased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 antagonists, and wherein homologues that have an increased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 agonists. The method can also include in step (b) comparing the binding affinity the homologue to the TALL-1 receptor or fragment of thereof to the binding affinity of wild-type TALL-1 and the TALL-1 receptor, and a step (d) of selecting homologues which have an increased binding affinity to the TALL-1 receptor or fragment of and a decreased TALL-1 biological activity.

Such methods can include cell-free or a cell-based assays. Binding assays and assays for detecting biological activity have been described above. The step of contacting can be performed by any suitable method. For example, cells expressing a TALL-1 receptor can be grown in liquid culture medium or grown on solid medium in which the liquid medium or the solid medium contains the compound to be tested in the presence or absence of a wild-type TALL-1. In addition, as described above, the liquid or solid medium contains components necessary for cell growth, such as assimilable carbon, nitrogen and micro-nutrients. In another embodiment, the TALL-1 protein, homologue, and/or the TALL-1 receptor and/or cell lysates containing such proteins can be immobilized on a substrate such

as: artificial membranes, organic supports, biopolymer supports and inorganic supports. The protein can be immobilized on the solid support by a variety of methods including adsorption, cross-linking (including covalent bonding), and entrapment. Adsorption can be through van der Waal's forces, hydrogen bonding, ionic bonding, or hydrophobic binding.

5 Exemplary solid supports for adsorption immobilization include polymeric adsorbents and ion-exchange resins. Solid supports can be in any suitable form, including in a bead form, plate form, or well form.

The present invention also provides the atomic coordinates that define the three dimensional structure of a sTALL-1, alone and in complex with eBCMA and with eBAFF-R.

10 First, the present inventors have determined the atomic coordinates that define the three dimensional structure of a crystalline TALL-1 (see Example 1 for details). Second, the present inventors have determined the atomic coordinates that define the three dimensional structure of a crystalline TALL-1 in complex with eBCMA (see Example 6 for details). Third, the present inventors have determined the atomic coordinates that define the three

15 dimensional structure of a crystalline TALL-1 in complex with eBAFF-R (see Example 6 for details). Using the guidance provided herein, one of skill in the art will be able to reproduce any of such structures and define atomic coordinates of such a structure.

As used herein, a "structure" of a protein refers to the components and the manner of arrangement of the components to constitute the protein. The "three dimensional

20 structure" or "tertiary structure" of the protein refers to the arrangement of the components of the protein in three dimensions. Such term is well known to those of skill in the art. It is also to be noted that the terms "tertiary" and "three dimensional" can be used interchangeably.

Example 1 describes the production of a sTALL-1, arranged in a crystalline manner

25 in a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217\text{\AA}$. The atomic coordinates determined from the crystal structure of TALL-1 are represented in Table 2. Table 2 represents the coordinates of a structure that has been refined to an R-factor of 23.6% (R_{free} -factor of 25.2%) against data extending to 3.0\AA resolution in space group $P6_322$, with ten sTALL-1 monomers in the asymmetric unit cell

30 having approximate dimensions of $a=b=234\text{\AA}$, $c=217\text{\AA}$. The atomic coordinates for the TALL-1 structure in Table 2 were deposited with the Protein Data Bank (PDB), operated by

the Research Collaboratory for Structural Bioinformatics (RCSB) (H.M.Berman, J.Westbrook, Z.Feng, G.Gilliland, T.N.Bhat, H.Weissig, I.N.Shindyalov, P.E.Bourne, The Protein Data Bank; *Nucleic Acids Research*, 28:235-242 (2000)), under PDB Deposit No. 1JH5 in June 2001, and such coordinates are incorporated herein by reference and such
5 coordinates are incorporated herein by reference.

Example 6 describes the production of a sTALL-1 complexed with eBCMA. The structure of sTALL-1 with eBCMA has been refined to an R-factor of 20.9% (R_{free} -factor of 23.4%) against data extending to 2.6 Å resolution in space group $P6_322$, with ten sTALL-1 monomers and seven entire and one partial eBCMA molecules in the asymmetric unit cell
10 having approximate dimensions of $a=b=234\text{Å}$, $c=217\text{Å}$ (Table 1B). Due to crystal packing, another two receptor binding sites were left unoccupied. The attached tables of atomic coordinates labeled Tables 3-12 define the coordinates of 10 structures of sTALL-1 determined from 10 complexes of sTALL-1 and BCMA, and the tables of atomic coordinates labeled 13-22 define the coordinates of 10 structures of the extracellular domains of BCMA
15 determined from the same complexes. Each structure of sTALL-1 is determined from the complex with one of the BCMA structures, such that Table 3 corresponds to Table 13, Table 4 corresponds to Table 14, and so on.. By way of example, the structure of sTALL-1 represented by the atomic coordinates in Table 3 was determined from a complex of sTALL-1 and BCMA, wherein the structure of the BCMA is represented by the atomic coordinates
20 in Table 13 (i.e., the structure represented by Table 3 was complexed with the structure represented by Table 13). Similarly, the structure of sTALL-1 represented by the atomic coordinates in Table 10 was determined from a complex of sTALL-1 and BCMA, wherein the structure of the BCMA is represented by the atomic coordinates in Table 20, and so on. The 10 different sTALL-1/BCMA complexes represented by the atomic coordinates provided
25 herein are representative of the complexes which together, form the unique virus-like assembly described for TALL-1/receptor (Table 3-Table 13; Table 4-Table 14; Table 5-Table 15; Table 6-Table 16; Table 7-Table 17; Table 8-Table 18; Table 9-Table 19; Table 10-Table 20; Table 11-Table 21; Table 12-Table 22).

Similar results are true for the sTALL-1 and eBAFF-R complex with a final
30 resolution of 3.0 Å (Table 1B). The attached tables of atomic coordinates labeled Tables 23-32 define the coordinates of 10 structures of the extracellular domains of eBAFF-R

determined from the complexes with sTALL-1. Together, the 10 different BAFF-R complexes represented by the atomic coordinates provided herein are representative of the structure of the receptor in the complex which together, form the unique virus-like assembly described for TALL-1/receptor.

5 One embodiment of the present invention includes a TALL-1, an eBMCA, or an eBAFF-R, in crystalline form. The present invention specifically exemplifies these crystalline forms. As used herein with regard to TALL-1 by way of example, the terms "crystalline TALL-1" and "TALL-1 crystal" both refer to crystallized TALL-1 and are intended to be used interchangeably. Preferably, a crystalline TALL-1 is produced using the
10 crystal formation method described herein, in particular according to the method disclosed in Example 1. A TALL-1 crystal of the present invention can comprise any crystal structure that comes from crystals formed in any of the allowable space groups for proteins (61 of them) and in one embodiment, crystallizes as an orthorhombic crystal lattice. In one aspect, a crystalline TALL-1 of the present invention includes TALL-1 molecules arranged in a
15 crystalline manner in a space group $P6_322$, with ten sTALL-1 monomers in the asymmetric unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217\text{\AA}$. According to the present invention, a unit cell having "approximate dimensions of" a given set of dimensions refers to a unit cell that has dimensions that are within plus (+) or minus (-) 2.0% of the specified unit cell dimensions. Such a small variation is within the scope of the invention since one
20 of skill in the art could obtain such variance by performing X-ray crystallography at different times on the same crystal. A preferred crystal of the present invention provides X-ray diffraction data for determination of atomic coordinates of the TALL-1 crystal to a resolution of about 4.0\AA , and preferably to about 3.2\AA , and preferably to about 3.0\AA .

One embodiment of the present invention includes a method for producing crystals
25 of TALL-1, TALL-1 in complex with its receptor, comprising combining the TALL-1 protein with a mother liquor and inducing crystal formation to produce the TALL-1 crystals. Although the production of crystals of TALL-1 is specifically described herein, it is to be understood that such processes as are described herein can be adapted by those of skill in the art to produce other crystals of TALL-1.

30 By way of example (i.e., this discussion applies to other crystals related to the invention, such as crystals of sTALL-1 and eBCMA or eBAFF-R), crystals of TALL-1 can

be formed using a solution containing TALL-1 in a mother liquor. A suitable mother liquor of the present invention comprises the solution used for crystallization as described in Example 1 that causes the protein to crystallize. There is some tolerance in the mother liquor conditions so that changes of up to 30% in buffer concentrations, pH units, and temperatures
5 can still yield crystals. Supersaturated solutions comprising TALL-1 can be induced to crystallize by several methods including, but not limited to, vapor diffusion, liquid diffusion, batch crystallization, constant temperature and temperature induction or a combination thereof. In one embodiment, supersaturated solutions of TALL-1 are induced to crystallize by hanging drop vapor diffusion. In a vapor diffusion method, TALL-1 molecule is
10 combined with a mother liquor as described above that will cause the protein solution to become supersaturated and form crystals at a constant temperature. Vapor diffusion is preferably performed under a controlled temperature and, by way of example, can be performed at 18° C. In a preferred embodiment, crystals are formed using the methods described in detail in the Examples section.

15 The crystalline TALL-1 of the present invention is analyzed by X-ray diffraction and, based on data collected from this procedure, models are constructed which represent the tertiary structure of the TALL-1 monomers. Therefore, one embodiment of the present invention includes a representation, or model, such as a computer model, of the three dimensional structure of TALL-1, as a monomer, trimer, cluster, or in complex with a
20 receptor. A computer model of the present invention can be produced using any suitable software modeling program, including, but not limited to, the graphical display program O (Jones et. al., *Acta Crystallography*, vol. A47, p. 110, 1991), the graphical display program GRASP, MOLSCRIPT 2.0 (Avatar Software AB, Heleneborgsgatan 21C, SE-11731 Stockholm, Sweden), the program CONTACTS from the CCP4 suite of programs (Bailey,
25 1994, *Acta Cryst.* D50:760-763), or the graphical display program INSIGHT. Suitable computer hardware useful for producing an image of the present invention are known to those of skill in the art (e.g., a Silicon Graphics Workstation).

A representation, or model, of the three dimensional structure of sTALL-1, eBCMA or eBAFF-R for which a crystal has been produced can also be determined using techniques
30 which include molecular replacement or SIR/MIR (single/multiple isomorphous replacement), or MAD (multiple wavelength anomalous diffraction) methods (Hendrickson

et al., 1997, *Methods Enzymol.*, 276:494-522). Methods of molecular replacement are generally known by those of skill in the art (generally described in Brunger, *Meth. Enzym.*, vol. 276, pp. 558-580, 1997; Navaza and Saludjian, *Meth. Enzym.*, vol. 276, pp. 581-594, 1997; Tong and Rossmann, *Meth. Enzym.*, vol. 276, pp. 594-611, 1997; and Bentley, *Meth. Enzym.*, vol. 276, pp. 611-619, 1997, each of which are incorporated by this reference herein in their entirety) and are performed in a software program including, for example, AmoRe (CCP4, *Acta Cryst. D*50, 760-763 (1994), SOLVE (Terwilliger et al., 1999, *Acta Crystallogr.*, D55:849-861), RESOLVE (Terwilliger, 2000, *Acta Crystallogr.*, D56:965-972) or XPLOR. Briefly, X-ray diffraction data is collected from the crystal of a crystallized target structure. The X-ray diffraction data is transformed to calculate a Patterson function. The Patterson function of the crystallized target structure is compared with a Patterson function calculated from a known structure (referred to herein as a search structure). The Patterson function of the crystallized target structure is rotated on the search structure Patterson function to determine the correct orientation of the crystallized target structure in the crystal. The translation function is then calculated to determine the location of the target structure with respect to the crystal axes. Once the crystallized target structure has been correctly positioned in the unit cell, initial phases for the experimental data can be calculated. These phases are necessary for calculation of an electron density map from which structural differences can be observed and for refinement of the structure. Preferably, the structural features (e.g., amino acid sequence, conserved di-sulphide bonds, and β -strands or β -sheets) of the search molecule are related to the crystallized target structure.

As used herein, the term "model" refers to a representation in a tangible medium of the three dimensional structure of a protein, polypeptide or peptide. For example, a model can be a representation of the three dimensional structure in an electronic file, on a computer screen, on a piece of paper (i.e., on a two dimensional medium), and/or as a ball-and-stick figure. Physical three-dimensional models are tangible and include, but are not limited to, stick models and space-filling models. The phrase "imaging the model on a computer screen" refers to the ability to express (or represent) and manipulate the model on a computer screen using appropriate computer hardware and software technology known to those skilled in the art. Such technology is available from a variety of sources including, for example, Evans and Sutherland, Salt Lake City, Utah, and Biosym Technologies, San Diego, CA. The

phrase "providing a picture of the model" refers to the ability to generate a "hard copy" of the model. Hard copies include both motion and still pictures. Computer screen images and pictures of the model can be visualized in a number of formats including space-filling representations, α carbon traces, ribbon diagrams and electron density maps. A variety of
5 such representations of the sTALL-1, eBCMA or eBAFF-R structural model are shown, for example, in the figures of the invention.

Preferably, a three dimensional structure of sTALL-1 provided by the present invention includes:

- (a) atomic coordinates determined by X-ray diffraction of a crystalline TALL-1;
- 10 (b) atomic coordinates selected from:
 - (1) atomic coordinates represented in any one of Tables 2-12;
 - (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7 Å over the backbone atoms in secondary structure elements of at least 50% of the residues
15 in a three dimensional structure represented by the atomic coordinates of (1); and
 - (3) atomic coordinates in any one of Tables 2-12 defining a portion of the TALL-1, wherein the portion of the TALL-1 comprises sufficient structural information to perform step (b); and
- (c) atomic coordinates defining the three dimensional structure of TALL-1 molecules
20 arranged in a crystalline manner in a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{Å}$, $c=217\text{Å}$.

Preferably, a three dimensional structure of BCMA or BAFF-R provided by the present invention includes:

- (a) atomic coordinates determined by X-ray diffraction of a crystalline BCMA or
25 crystalline BAFF-R;
- (b) atomic coordinates selected from:
 - (1) atomic coordinates represented in any one of Tables 13-33;
 - (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7 Å over the backbone atoms in secondary structure elements of at least 50% of the residues
30 in a three dimensional structure represented by said atomic coordinates of (1); and

(3) atomic coordinates in any one of Tables 13-22 defining a portion of said BCMA, wherein the portion of said BCMA comprises sufficient structural information to perform step (b);

(4) atomic coordinates in any one of Tables 14-33 defining a portion of said BAFF-R, wherein the portion of said BAFF-R comprises sufficient structural information to perform step (b);

(c) atomic coordinates defining the three dimensional structure of BCMA molecules or BAFF-R molecules arranged in a crystalline manner in a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217$.

In one aspect as described above, a three dimensional structure of sTALL-1, eBCMA or eBAFF-R provided by the present invention includes a structure wherein the structure has an average root-mean-square deviation (RMSD) of equal to or less than about 1.7\AA over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by the atomic coordinates of any one of the referenced tables of atomic coordinates. Such a structure can be referred to as a structural homologue of the sTALL-1, eBCMA or eBAFF-R structures defined by one of the corresponding referenced tables of atomic coordinates. Preferably, the structure has an average root-mean-square deviation (RMSD) of equal to or less than about 1.6\AA over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by the atomic coordinates of any one of the corresponding reference tables, or equal to or less than about 1.5\AA , or equal to or less than about 1.4\AA , or equal to or less than about 1.3\AA , or equal to or less than about 1.2\AA , or equal to or less than about 1.1\AA , or equal to or less than about 1.0\AA , or equal to or less than about 0.9\AA , or equal to or less than about 0.8\AA , or equal to or less than about 0.7\AA , or equal to or less than about 0.6\AA , or equal to or less than about 0.5\AA , or equal to or less than about 0.4\AA , or equal to or less than about 0.3\AA , or equal to or less than about 0.2\AA , over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by the atomic coordinates of any one of the corresponding reference tables of atomic coordinates. In another aspect, a three dimensional structure of a sTALL-1, eBCMA or eBAFF-R provided by the present invention includes a structure wherein the structure has the recited RMSD over the backbone atoms in secondary structure elements of at least 75% of the

residues in a three dimensional structure represented by the atomic coordinates of any one of the corresponding referenced tables of atomic coordinates, and more preferably at least about 80%, and more preferably at least about 85%, and more preferably at least about 90%, and more preferably at least about 95%, and most preferably, about 100% of the residues in
5 a three dimensional structure represented by the atomic coordinates of any one of the corresponding referenced tables.

In one embodiment, the RMSD of a structural homologue of sTALL-1, eBCMA or eBAFF-R can be extended to include atoms of amino acid side chains. As used herein, the phrase "common amino acid side chains" refers to amino acid side chains that are common
10 to both the structural homologue and to the structure that is actually represented by such atomic coordinates (e.g., a structure represented by one of the tables of atomic coordinates). Preferably, at least 50% of the structure has an average root-mean-square deviation (RMSD) from common amino acid side chains in a three dimensional structure represented by the atomic coordinates of one of the referenced tables of equal to or less than about 1.7 Å, or
15 equal to or less than about 1.6 Å, equal to or less than about 1.5 Å, or equal to or less than about 1.4 Å, or equal to or less than about 1.3 Å, or equal to or less than about 1.2 Å, or equal to or less than about 1.1 Å, or equal to or less than about 1.0 Å, or equal to or less than about 0.9 Å, or equal to or less than about 0.8 Å, or equal to or less than about 0.7 Å, or equal to or less than about 0.6 Å, or equal to or less than about 0.5 Å, or equal to or less than about
20 0.4 Å, or equal to or less than about 0.3 Å, or equal to or less than about 0.2 Å. In another embodiment, a three dimensional structure of sTALL-1, eBCMA or eBAFF-R provided by the present invention includes a structure wherein at least about 75% of such structure has the recited average root-mean-square deviation (RMSD) value, and more preferably, at least about 85% of such structure has the recited average root-mean-square deviation (RMSD)
25 value, and most preferably, about 95% of such structure has the recited average root-mean-square deviation (RMSD) value.

In addition to having the recited RMSD values, a structural homologue of sTALL-1, eBCMA or eBAFF-R should additionally meet the criteria for amino acid sequence identity which was discussed in detail previously herein. For a given amino acid sequence or amino
30 acid residue to correspond to an amino acid region or amino acid position in another sequence, the position of the sequence or residue in the query sequence should align to the

position of the region or residue in the compared sequence using a standard alignment program in the art, but particularly, using the programs BLOCKS (GIBBS) and/or MAST (Henikoff et al., 1995, *Gene*, 163, 17-26; Henikoff et al., 1994, *Genomics*, 19, 97-107), using standard manufacturer defaults.

5 Another structure that is useful in the methods of the present invention is a structure that is defined by the atomic coordinates in any one of the referenced tables of atomic coordinates that define a portion of the sTALL-1, eBCMA or eBAFF-R, wherein the portion of the sTALL-1, eBCMA or eBAFF-R comprises sufficient structural information to perform structure based drug design (described below). Suitable portions of sTALL-1, eBCMA or
10 eBAFF-R that could be modeled and used in structure based drug design will be apparent to those of skill in the art. The present inventors have also identified multiple sites of interest based on the structure of sTALL-1, eBCMA or eBAFF-R (described in detail above and in the Examples). Structures comprising these portions (e.g., a receptor or ligand binding region) would be encompassed by the present invention.

15 Accordingly, another embodiment of the present invention relates to a method of structure-based identification of compounds which potentially bind to TALL-1. The method includes the steps of: (a) obtaining atomic coordinates that define the three dimensional structure of TALL-1 (described below); and (b) selecting candidate compounds for binding to the TALL-1 by performing structure based drug design with the structure of (a), wherein
20 the step of selecting is performed in conjunction with computer modeling. The atomic coordinates used in the method are selected from:

(i) atomic coordinates determined by X-ray diffraction of a crystalline TALL-

1;

(ii) atomic coordinates selected from:

25 (1) atomic coordinates represented in any one of Tables 2-12;

 (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7 Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by the atomic
30 coordinates of (1); and

(3) atomic coordinates in any one of Tables 2-12 defining a portion of the TALL-1, wherein the portion of the TALL-1 comprises sufficient structural information to perform step (b); and

(iii) atomic coordinates defining the three dimensional structure of TALL-1 molecules arranged in a crystalline manner in a space group P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217 Å.

In one embodiment, the method further comprises a step (c) of selecting candidate compounds of (b) that inhibit the biological activity of TALL-1. In one aspect, this step of selecting comprises (i) contacting the candidate compound identified in step (b) with TALL-1; and (ii) measuring the biological activity of the TALL-1, as compared to in the absence of the candidate compound.

In another embodiment, the method further comprises a step (c) of selecting candidate compounds of (b) that inhibit the binding of TALL-1 to a TALL-1 receptor. In this aspect, the step (c) can include: (i) contacting the candidate compound identified in step (b) with the TALL-1 or a fragment thereof and a TALL-1 receptor or TALL-1 receptor binding fragment thereof under conditions in which a TALL-1-TALL-1 receptor complex can form in the absence of the candidate compound; and (ii) measuring the binding of the TALL-1 or fragment thereof to bind to the TALL-1 receptor or fragment thereof, wherein a candidate inhibitor compound is selected when there is a decrease in the binding of the TALL-1 or fragment thereof to the TALL-1 receptor or fragment thereof, as compared to in the absence of the candidate inhibitor compound. A TALL-1 receptor used in these embodiments can include BCMA, BAFF-R and TACI.

In the general method, the step (b) of selecting can include identifying candidate compounds for binding to a receptor binding site of the TALL-1 protein, the receptor binding site comprising an amino acid residue selected from the group consisting of Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222. In another aspect, step (b) of selecting comprises identifying candidate compounds for binding to the TALL-1 such that trimer-trimer interactions between trimers of TALL-1 monomers is inhibited. For example, the step of selecting can include identifying candidate compounds for binding to TALL-1 at a site including an amino acid residue selected from the group consisting of: Gln144, Ile150, Leu169, Phe172, Tyr192, Phe194,

Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285.

Another embodiment of the present invention relates to a method to construct a three dimensional model of TALL-1 protein or homologue thereof. The method includes the steps
5 of: (a) obtaining atomic coordinates that define the three dimensional structure of TALL-1; and (b) performing computer modeling with the atomic coordinates of (a) and to construct a model of a three dimensional structure of a TALL-1 or homologue thereof. the atomic coordinates are the same as those described above for the method of identifying compounds that bind to TALL-1.

10 Yet another embodiment of the present invention relates to a method of structure-based identification of compounds which potentially bind to a TALL-1 receptor selected from the group consisting of BCMA and BAFF-R, comprising: (a) obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R; and (b) selecting candidate compounds for binding to the BCMA or BAFF-R by performing structure
15 based drug design with the structure of (a), wherein the step of selecting is performed in conjunction with computer modeling. The atomic coordinates used in the method are selected from:

(a) atomic coordinates determined by X-ray diffraction of a crystalline BCMA or crystalline BAFF-R;

20 (b) atomic coordinates selected from the group consisting of:

(1) atomic coordinates represented in any one of Tables 13-33;

(2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7 Å over the backbone atoms in secondary structure elements of at least 50%
25 of the residues in a three dimensional structure represented by the atomic coordinates of (1);

(3) atomic coordinates in any one of Tables 13-22 defining a portion of the BCMA, wherein the portion of the BCMA comprises sufficient structural information to perform step (b); and

(4) atomic coordinates in any one of Tables 14-33 defining a portion of the BAFF-R, wherein the portion of the BAFF-R comprises sufficient structural information to perform step (b); and

(c) atomic coordinates defining the three dimensional structure of BCMA molecules or BAFF-R molecules arranged in a crystalline manner in a space group P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217.

Yet another embodiment of the present invention relates to a method to construct a three dimensional model of BCMA, BAFF-R, TACI, or a homologue thereof, comprising: (a) obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R (as described for the method above); and, (b) performing computer modeling with the atomic coordinates of (a) and an amino acid sequence corresponding to BCMA, BAFF-R or TACI to construct a model of a three dimensional structure of the BCMA, BAFF-R or TACI, or homologue thereof.

The structures and atomic coordinates used to perform the above-described method have been described in detail above and in the Examples section, and include any structural homologues of TALL-1 described herein or in other embodiments of drug design described below, any structural homologues of a TALL-1 receptor described herein. In general, terms and definitions used to describe this embodiment related to the method of structure-based identification of compounds which potentially bind to TALL-1 will apply to the other methods of structure-based identification of compounds that bind to a TALL-1 receptor, or to methods of designing compounds that mimic the structure of TALL-1 or TALL-1 receptors.

According to the present invention, the phrase "obtaining atomic coordinates that define the three dimensional structure of TALL-1" is defined as any means of obtaining providing, supplying, accessing, displaying, retrieving, or otherwise making available the atomic coordinates defining any three dimensional structure of TALL-1 as described herein. For example, the step of obtaining can include, but is not limited to, accessing the atomic coordinates for the structure from a database or other source; importing the atomic coordinates for the structure into a computer or other database; displaying the atomic coordinates and/or a model of the structure in any manner, such as on a computer, on paper,

etc.; and determining the three dimensional structure of TALL-1 described by the present invention *de novo* using the guidance provided herein.

The second step of the method of structure based identification of compounds of the present invention includes selecting a candidate compound for binding to and/or inhibiting the biological activity of the TALL-1 represented by the structure model by performing
5 structure based drug design with the model of the structure. According to the present invention, the step of "selecting" can refer to any screening process, modeling process, design process, or other process by which a compound can be selected as useful for binding to and enhancing or inhibiting the activity of TALL-1 according to the present invention. Methods
10 of structure based identification of compounds are described in detail below. As discussed above, TALL-1 is involved in the regulation of B cell development, proliferation and survival, and therefore, the selection of compounds that compete with, disrupt or otherwise inhibit the biological activity of TALL-1 (or its receptors) are highly desirable. In addition, in some embodiments, compounds which enhance, increase, or otherwise agonize the
15 biological activity of TALL-1 (or its receptors) may be desirable, such as in immunodeficiency diseases or vaccine administration. Any of such compounds can be designed using structure based drug design using models of the structures disclosed herein. Until the discovery of the three dimensional structure of the present invention, the only information available for the development of therapeutic compounds based on TALL-1 was
20 based on the primary sequence of TALL-1 and perhaps, mutagenesis studies directed to the isolated protein.

Structure based identification of compounds (e.g., structure based drug design, structure based compound screening, or structure based structure modeling) refers to the prediction or design of a conformation of a peptide, polypeptide, protein (e.g., TALL-1), or
25 to the prediction or design of a conformational interaction between such protein, peptide or polypeptide, and a candidate compound, by using the three dimensional structure of the peptide, polypeptide or protein. Typically, structure based identification of compounds is performed with a computer (e.g., computer-assisted drug design, screening or modeling). For example, generally, for a protein to effectively interact with (e.g., bind to) a compound, it is
30 necessary that the three dimensional structure of the compound assume a compatible conformation that allows the compound to bind to the protein in such a manner that a desired

result is obtained upon binding. Knowledge of the three dimensional structure of TALL-1 and two of its cognate receptors enables a skilled artisan to design a compound having such compatible conformation, or to select such a compound from available libraries of compounds and/or structures thereof. For example, knowledge of the three dimensional structure of the receptor binding site of TALL-1 or the ligand binding site of a TALL-1 receptor enables one of skill in the art to design or select a compound structure that is predicted to bind to the TALL-1 or its receptor at that site and result in, for example, inhibition of the binding of a wild-type TALL-1 to the receptor, thereby inhibiting a biological response such as activation of B cell proliferation or maturation. In addition, for example, knowledge of the three dimensional structure of TALL-1 enables a skilled artisan to design an analog (structural homologue) of TALL-1 or an analog of TALL-1 receptor.

Suitable structures and models useful for structure based drug design are disclosed herein. Preferred target structures to use in a method of structure based drug design include any representations of structures produced by any modeling method disclosed herein, including molecular replacement and fold recognition related methods.

According to the present invention, the step of selecting or designing a compound for testing in a method of structure based identification of the present invention can include creating a new chemical compound structure or searching databases of libraries of known compounds (e.g., a compound listed in a computational screening database containing three dimensional structures of known compounds). Designing can also be performed by simulating chemical compounds having substitute moieties at certain structural features. The step of designing can include selecting a chemical compound based on a known function of the compound. A preferred step of designing comprises computational screening of one or more databases of compounds in which the three dimensional structure of the compound is known and is interacted (e.g., docked, aligned, matched, interfaced) with the three dimensional structure of a TALL-1 protein or receptor by computer (e.g. as described by Humblet and Dunbar, *Animal Reports in Medicinal Chemistry*, vol. 28, pp. 275-283, 1993, M Venuti, ed., Academic Press). The compound itself, if identified as a suitable candidate by the method of the invention, can be synthesized and tested directly with the TALL-1 protein or a TALL-1 receptor in a biological assay. Methods to synthesize suitable chemical compounds are known to those of skill in the art and depend upon the structure of the

chemical being synthesized. Methods to evaluate the bioactivity of the synthesized compound depend upon the bioactivity of the compound (e.g., inhibitory or stimulatory) and are discussed herein.

Various other methods of structure-based drug design are disclosed in Maulik et al.,
5 1997, *Molecular Biotechnology: Therapeutic Applications and Strategies*, Wiley-Liss, Inc.,
which is incorporated herein by reference in its entirety. Maulik et al. disclose, for example,
methods of directed design, in which the user directs the process of creating novel molecules
from a fragment library of appropriately selected fragments; random design, in which the user
uses a genetic or other algorithm to randomly mutate fragments and their combinations while
10 simultaneously applying a selection criterion to evaluate the fitness of candidate ligands; and
a grid-based approach in which the user calculates the interaction energy between three
dimensional receptor structures and small fragment probes, followed by linking together of
favorable probe sites.

In a molecular diversity strategy, large compound libraries are synthesized, for
15 example, from peptides, oligonucleotides, carbohydrates and/or synthetic organic molecules,
using biological, enzymatic and/or chemical approaches. The critical parameters in
developing a molecular diversity strategy include subunit diversity, molecular size, and
library diversity. The general goal of screening such libraries is to utilize sequential
application of combinatorial selection to obtain high-affinity ligands for a desired target, and
20 then to optimize the lead molecules by either random or directed design strategies. Methods
of molecular diversity are described in detail in Maulik, et al., *ibid*.

Maulik et al. also disclose, for example, methods of directed design, in which the user
directs the process of creating novel molecules from a fragment library of appropriately
selected fragments; random design, in which the user uses a genetic or other algorithm to
25 randomly mutate fragments and their combinations while simultaneously applying a selection
criterion to evaluate the fitness of candidate ligands; and a grid-based approach in which the
user calculates the interaction energy between three dimensional receptor structures and small
fragment probes, followed by linking together of favorable probe sites.

In the present method of structure based identification of compounds, it is not
30 necessary to align the structure of a candidate chemical compound (i.e., a chemical
compound being analyzed in, for example, a computational screening method of the present

invention) to each residue in a target site (target sites will be discussed in detail below). Suitable candidate chemical compounds can align to a subset of residues described for a target site. Preferably, a candidate chemical compound comprises a conformation that promotes the formation of covalent or noncovalent crosslinking between the target site and the candidate chemical compound. In one aspect, a candidate chemical compound binds to a surface adjacent to a target site to provide an additional site of interaction in a complex. When designing an antagonist (i.e., a chemical compound that inhibits the biological activity of an TALL-1), for example, the antagonist should bind with sufficient affinity to the target binding site or substantially prohibit a ligand (e.g., a molecule that specifically binds to the target site) from binding to a target site. It will be appreciated by one of skill in the art that it is not necessary that the complementarity between a candidate chemical compound and a target site extend over all residues specified here in order to inhibit or promote binding of a ligand.

In general, the design of a chemical compound possessing stereochemical complementarity can be accomplished by techniques that optimize, chemically or geometrically, the "fit" between a chemical compound and a target site. Such techniques are disclosed by, for example, Sheridan and Venkataraghavan, *Acc. Chem Res.*, vol. 20, p. 322, 1987; Goodford, *J. Med. Chem.*, vol. 27, p. 557, 1984; Beddell, *Chem. Soc. Reviews*, vol. 279, 1985; Hol, *Angew. Chem.*, vol. 25, p. 767, 1986; and Verlinde and Hol, *Structure*, vol. 2, p. 577, 1994, each of which are incorporated by this reference herein in their entirety.

One embodiment of the present invention for structure based drug design comprises identifying a chemical compound that complements the shape of TALL-1 or a TALL-1 receptor, including a portion of TALL-1 or a TALL-1 receptor. Such method is referred to herein as a "geometric approach". In a geometric approach, the number of internal degrees of freedom (and the corresponding local minima in the molecular conformation space) is reduced by considering only the geometric (hard-sphere) interactions of two rigid bodies, where one body (the active site) contains "pockets" or "grooves" that form binding sites for the second body (the complementing molecule, such as a ligand).

The geometric approach is described by Kuntz et al., *J. Mol. Biol.*, vol. 161, p. 269, 1982, which is incorporated by this reference herein in its entirety. The algorithm for chemical compound design can be implemented using the software program DOCK Package,

Version 1.0 (available from the Regents of the University of California). Pursuant to the Kuntz algorithm, the shape of the cavity or groove on the surface of a structure (e.g., TALL-1) at a binding site or interface is defined as a series of overlapping spheres of different radii. One or more extant databases of crystallographic data (e.g., the Cambridge Structural Database System maintained by University Chemical Laboratory, Cambridge University, 5 Lensfield Road, Cambridge CB2 1EW, U.K.) or the Protein Data Bank maintained by Brookhaven National Laboratory, is then searched for chemical compounds that approximate the shape thus defined.

Chemical compounds identified by the geometric approach can be modified to satisfy 10 criteria associated with chemical complementarity, such as hydrogen bonding, ionic interactions or Van der Waals interactions.

Another embodiment of the present invention for structure based identification of compounds comprises determining the interaction of chemical groups ("probes") with an active site at sample positions within and around a binding site or interface, resulting in an 15 array of energy values from which three dimensional contour surfaces at selected energy levels can be generated. This method is referred to herein as a "chemical-probe approach." The chemical-probe approach to the design of a chemical compound of the present invention is described by, for example, Goodford, *J. Med. Chem.*, vol. 28, p. 849, 1985, which is incorporated by this reference herein in its entirety, and is implemented using an appropriate 20 software package, including for example, GRID (available from Molecular Discovery Ltd., Oxford OX2 9LL, U.K.). The chemical prerequisites for a site-complementing molecule can be identified at the outset, by probing the active site of a TALL-1 protein, for example, (e.g., as represented by the atomic coordinates shown in one of the tables herein) with different chemical probes, e.g., water, a methyl group, an amine nitrogen, a carboxyl oxygen and/or 25 a hydroxyl. Preferred sites for interaction between an active site and a probe are determined. Putative complementary chemical compounds can be generated using the resulting three dimensional pattern of such sites.

According to the present invention, suitable candidate compounds to test using the method of the present invention include proteins, peptides or other organic molecules, and 30 inorganic molecules. Suitable organic molecules include small organic molecules. Peptides refer to small molecular weight compounds yielding two or more amino acids upon

hydrolysis. A polypeptide is comprised of two or more peptides. As used herein, a protein is comprised of one or more polypeptides. Preferred therapeutic compounds to design include peptides composed of "L" and/or "D" amino acids that are configured as normal or retroinverso peptides, peptidomimetic compounds, small organic molecules, or homo- or hetero-polymers thereof, in linear or branched configurations.

In one embodiment, the compound that is identified by the method of the present invention originates from a compound having chemical and/or stereochemical complementarity with a site on a TALL-1 protein or a TALL-1 receptor. Such complementarity is characteristic of a compound that matches the surface of the enzyme either in shape or in distribution of chemical groups and binds to TALL-1 or a TALL-1 receptor to inhibit binding of TALL-1 to its receptor, for example, or to otherwise inhibit the biological activity of TALL-1 or its receptor and/or inhibit B cell proliferation, maturation, development or survival in a cell expressing TALL-1 receptor. More preferably, a compound that binds to a receptor binding site on TALL-1 or to a ligand binding site on a TALL-1 receptor associates with an affinity of at least about 10^{-6} M, and more preferably with an affinity of at least about 10^{-7} M, and more preferably with an affinity of at least about 10^{-8} M.

The general sites of both TALL-1 and its receptors as targets for structure based drug design or identification of candidate compounds and lead compounds (i.e., target sites), have been discussed in detail above, although other sites may become apparent to those of skill in the art using the structures provided herein. The sites generally include for TALL-1 the sites involved in trimer formation, trimer-trimer interactions (including the flap region) and receptor binding sites as described in detail herein. For TALL-1 receptors, the sites include the ligand binding sites. Combinations of any of these general sites are also suitable target sites. Even if some of such sites were generally known or hypothesized to be important sites prior to the present invention based on the linear sequence and mutational analysis or binding studies of TALL-1, the present invention actually defines the sites in three dimensions and confirms or newly identifies residues that are important targets that could not be confirmed or identified prior to the present invention. The use of any of these target sites or any other sites that can be elucidated as a result of the determination of the three dimensional structure described herein is novel and encompassed by the present invention. Many of these target sites are further described and illustrated in the Figures and Examples of the invention.

The Examples section provides specific detail regarding the structure of TALL-1 and its receptors BCMA and BAFF-R and target sites of TALL-1, BCMA and BAFF-R based on the three-dimensional structures described herein, including the identification of important residues in the structures. It is to be understood, however, that one of skill in the art, using the description of these specific structures provided herein, will be able to identify compounds that are potential candidates for modulating the biological activity of TALL-1, APRIL and the receptors for TALL-1 and APRIL, including TACI. All such embodiments are encompassed by the present invention.

A candidate compound for binding to or otherwise modulating the activity of TALL-1 or its receptors, including to one of the preferred target sites described above, is identified by one or more of the methods of structure-based identification discussed above. As used herein, a "candidate compound" refers to a compound that is selected by a method of structure-based identification described herein as having a potential for binding to TALL-1 or its receptors on the basis of a predicted conformational interaction between the candidate compound and the target site of TALL-1 or its receptors. The ability of the candidate compound to actually bind to TALL-1 or its receptors can be determined using techniques known in the art, as discussed in some detail below. A "putative compound" is a compound with an unknown regulatory activity, at least with respect to the ability of such a compound to bind to and/or regulate TALL-1 or its receptors as described herein. Therefore, a library of putative compounds can be screened using structure based identification methods as discussed herein, and from the putative compounds, one or more candidate compounds for binding to or mimicking the target TALL-1 or its receptor can be identified. Alternatively, a candidate compound for binding to or mimicking TALL-1 or its receptors can be designed *de novo* using structure based drug design, also as discussed above.

Accordingly, in one aspect of the present invention, the method of structure-based identification of compounds that potentially bind to or modulate (regulate) the activity of an TALL-1 or its receptors further includes steps which confirm whether or not a candidate compound has the predicted properties with respect to its effect on the actual TALL-1 or receptor. In one embodiment, the candidate compound is predicted to be an inhibitor of the binding of TALL-1 to at least one of its receptors, and the method further includes producing or otherwise obtaining a candidate compound selected in the structure based method and

determining whether the compound actually has the predicted effect on the TALL-1 protein or its biological activity. For example, one can additionally contact the candidate compound selected in the structure based identification method with TALL-1 or a fragment thereof under conditions in which the TALL-1 binds to its receptor in the absence of the candidate compound; and measuring the binding affinity of TALL-1 or fragment thereof for its receptor
5 or a fragment thereof. In this example (binding), a candidate inhibitor compound is selected as a compound that inhibits the binding of TALL-1 to its receptor when there is a decrease in the binding affinity of TALL-1 or fragment thereof for the substrate or fragment thereof, as compared to in the absence of the candidate inhibitor compound. This experiment can be
10 used to identify compounds that inhibit the binding of TALL-1 to its receptor via binding to TALL-1 or its receptor by simple manipulation of the order of adding the compounds to the assay, for example.

In another embodiment, the candidate compound is predicted to inhibit the biological activity of TALL-1 or its receptor, and the method further comprises contacting the actual
15 candidate compound selected by the structure-based identification method with TALL-1 or its receptor or a targeted fragment thereof, under conditions wherein in the absence of the compound, TALL-1 and/or its receptor are biologically active, and measuring the ability of the candidate compound to inhibit the activity of TALL-1 or its receptor.

In another embodiment, the candidate compound, or modeled TALL-1 or TALL-1
20 receptor structure in some embodiments (described below), is predicted to be a mimic or homologue of a natural TALL-1 or its receptor, respectively, and is predicted to have modified biological activity as compared to the natural TALL-1 or its receptor. For example, one can model and then produce and test a TALL-1 homologue that has different receptor binding affinity as compared to the natural TALL-1, or a homologue that increased or
25 decreased biological activity as compared to the natural TALL-1. Such homologues can be useful in various biological assays or as competitive inhibitors.

In one embodiment, the conditions under which TALL-1 or a TALL-1 receptor according to the present invention is contacted with a candidate compound, such as by mixing, are conditions in which the protein is not stimulated (activated) or bound to a natural
30 ligand or receptor if essentially no candidate compound is present. In one aspect, a natural ligand or substrate can be added after contact with the candidate compound to determine the

effect of the compound on the biological activity of TALL-1 or its receptor. Alternatively, this aspect can be designed simply to determine whether the candidate compound binds to the TALL-1 or its receptor (i.e., in the absence of any additional testing, such as by addition of the receptor or ligand, respectively). For example, such conditions include normal culture
5 conditions in the absence of a stimulatory compound or binding ligand or receptor.

In another embodiment, the conditions under which TALL-1 or its receptor according to the present invention is contacted with a candidate compound, such as by mixing, are conditions in which the protein is normally bound by a ligand or receptor, or activated, if essentially no candidate compound is present. Such conditions can include, for example,
10 contact of TALL-1 or its receptor with the appropriate binding ligands or other stimulatory molecule. In this embodiment, the candidate compound can be contacted with TALL-1 or its receptor prior to the contact of TALL-1 or its receptor with the binding ligand (e.g., to determine whether the candidate compound blocks or otherwise inhibits the binding of TALL-1 or its receptor to its ligand, or inhibits the biological activity of TALL-1 or its
15 receptor), or after contact of TALL-1 or its receptor with the binding ligand (e.g., to determine whether the candidate compound downregulates, or reduces the biological activity of TALL-1 or its receptor after the initial contact with the binding ligand).

The present methods involve contacting TALL-1 or its receptor with the candidate compound being tested for a sufficient time to allow for binding to, activation or inhibition
20 of the enzyme by the candidate compound. The period of contact with the candidate compound being tested can be varied depending on the result being measured, and can be determined by one of skill in the art. For example, for binding assays, a shorter time of contact with the candidate compound being tested is typically suitable, than when activation is assessed. As used herein, the term "contact period" refers to the time period during which
25 TALL-1 or its receptor is in contact with the compound being tested. The term "incubation period" refers to the entire time during which cells expressing TALL-1 or its receptor, for example, are allowed to grow or incubate prior to evaluation, and can be inclusive of the contact period. Thus, the incubation period includes all of the contact period and may include a further time period during which the compound being tested is not present but
30 during which growth or cellular events are continuing (in the case of a cell based assay) prior

to scoring. It will be recognized that shorter incubation times are preferable because compounds can be more rapidly screened.

In accordance with the present invention, a cell-based assay is conducted under conditions that are effective to screen candidate compounds selected in the structure-based
5 identification method to confirm whether such compounds are useful as predicted. Effective conditions include, but are not limited to, appropriate media, temperature, pH and oxygen conditions that permit the growth of the cell that expresses TALL-1 or its receptor. An appropriate, or effective, medium refers to any medium in which a cell that naturally or recombinantly expresses TALL-1 or its receptor, when cultured, is capable of cell growth and
10 expression of TALL-1 or its receptor. Such a medium is typically a solid or liquid medium comprising growth factors and assimilable carbon, nitrogen, sulfur and phosphate sources, as well as appropriate salts, minerals, metals and other nutrients, such as vitamins. Culturing is carried out at a temperature, pH and oxygen content appropriate for the cell. Such culturing conditions are within the expertise of one of ordinary skill in the art.

15 Cells that are useful in the cell-based assays of the present invention include any cell that expresses TALL-1 or its receptor and particularly, other components related to B cell development, proliferation, maturation or survival.

The assay of the present invention can also be a non-cell based assay. In this embodiment, the candidate compound can be directly contacted with an isolated TALL-1 or
20 its receptor, or a portion thereof (e.g., a portion comprising a receptor or ligand binding region), and the ability of the candidate compound to bind to the protein or portion thereof can be evaluated. The assay can, if desired, additionally include the step of further analyzing whether candidate compounds which bind to TALL-1 or its receptor are capable of increasing or decreasing the activity of TALL-1 or its receptor. Such further steps can be performed by
25 cell-based assay, as described above, or by a non-cell-based assay that measures a parameter of TALL-1 or its receptor activity. For example, TALL-1 or its receptor can be immobilized on a solid support and evaluated for binding to a candidate compound and additionally, activity can be measured if the appropriate conditions and substrates are provided. Proteins can be immobilized on a substrate such as: artificial membranes, organic supports,
30 biopolymer supports and inorganic supports. The protein can be immobilized on the solid support by a variety of methods including adsorption, cross-linking (including covalent

bonding), and entrapment. Adsorption can be through van der Waal's forces, hydrogen bonding, ionic bonding, or hydrophobic binding. Exemplary solid supports for adsorption immobilization include polymeric adsorbents and ion-exchange resins. Solid supports can be in any suitable form, including in a bead form, plate form, or well form.

5 In one embodiment, a BIAcore machine can be used to determine the binding constant of a complex between TALL-1 or its receptor and a candidate compound or between TALL-1 and its receptor, for example, in the presence and absence of the candidate compound. The dissociation constant for the complex can be determined by monitoring changes in the refractive index with respect to time as buffer is passed over the chip
10 (O'Shannessy et al. *Anal. Biochem.* 212:457-468 (1993); Schuster et al., *Nature* 365:343-347 (1993)). Contacting a candidate compound at various concentrations with TALL-1 or its receptor and monitoring the response function (e.g., the change in the refractive index with respect to time) allows the complex dissociation constant to be determined in the presence of the candidate compound.

15 Other suitable assays for measuring the binding of a candidate compound to TALL-1 or its receptor and/or for measuring the ability of such compound to affect the binding of an TALL-1 to its receptor include, for example, immunoassays such as enzyme linked immunoabsorbent assays (ELISA) and radioimmunoassays (RIA), or determination of binding by monitoring the change in the spectroscopic or optical properties of the TALL-1
20 or its receptor, through fluorescence, UV absorption, circular dichroism, or nuclear magnetic resonance (NMR).

Candidate compounds identified by the present invention can include agonists of TALL-1 or TALL-1 receptor activity and antagonists of TALL-1 or TALL-1 receptor activity, with the identification of antagonists or inhibitors being preferred.

25 Yet another embodiment of the present invention relates to a method to produce a TALL-1 or TALL-1 receptor homologue with modified biological activity as compared to a natural TALL-1 or TALL-1 receptor. This method includes the steps of: (a) obtaining atomic coordinates that define the three dimensional structure of TALL-1 or a TALL-1 receptor, including any of the TALL-1 or TALL-1 receptor three dimensional structures or
30 atomic coordinates described herein; (b) using computer modeling of the atomic coordinates in (a) to identify at least one site in the TALL-1 or TALL-1 receptor structure that is

predicted to contribute to the biological activity of TALL-1 or the TALL-1 receptor; and (c) modifying the at least one site in the TALL-1 or TALL-1 receptor protein to produce a TALL-1 homologue or TALL-1 receptor homologue which is predicted to have modified biological activity as compared to a natural TALL-1 or TALL-1 receptor. The final step of
5 modifying the site on TALL-1 or a TALL-1 receptor can be performed by producing a "virtual TALL-1 or TALL-1 receptor homologue" on a computer, such as by generating a computer model of a TALL-1 or TALL-1 receptor homologue, or by modifying a TALL-1 or TALL-1 receptor to produce the homologue, such as by classical mutagenesis or recombinant technology.

10 The atomic coordinates that define the three dimensional structure of TALL-1 or a TALL-1 receptor and the step of obtaining such coordinates have been described in detail previously herein with regard to the method of structure based identification of compounds. Computer modeling methods suitable for modeling the atomic coordinates to identify sites
15 in a TALL-1 or TALL-1 receptor structure that are predicted to contribute to the biological activity of a TALL-1 or TALL-1 receptor, as well as for modeling homologues of TALL-1 or a TALL-1 receptor, have been discussed generally above. A variety of computer software programs for modeling and analyzing three dimensional structures of proteins are publicly available. The Examples section describes in detail the use of a few of such programs. Such computer software programs include, but are not limited to, the graphical display program
20 O (Jones et. al., *Acta Crystallography*, vol. A47, p. 110, 1991), the graphical display program GRASP, MOLSCRIPT 2.0 (Avatar Software AB, Heleneborgsgatan 21C, SE-11731 Stockholm, Sweden), the program CONTACTS from the CCP4 suite of programs (Bailey, 1994, *Acta Cryst.* D50:760-763), or the graphical display program INSIGHT.

Once target sites for modification on a TALL-1 or TALL-1 receptor are identified,
25 TALL-1 or TALL-1 receptor homologues having modifications at these sites can be produced and evaluated to determine the effect of such modifications on TALL-1 or TALL-1 receptor biological activity. In one embodiment, a TALL-1 or TALL-1 receptor homologue can be modeled on a computer to produce a computer model of a TALL-1 or TALL-1 receptor homologue which predicts the effects of given modifications on the structure of the protein
30 and its subsequent interaction with other molecules. Such computer modeling techniques are well known in the art.

In another aspect, or subsequent to an initial computer generation and evaluation of a TALL-1 or TALL-1 receptor homologue model, an actual TALL-1 or TALL-1 receptor homologue can be produced and evaluated by modifying target sites of a natural TALL-1 or TALL-1 receptor to produce a modified or mutant TALL-1 or TALL-1 receptor (described
5 in detail above).

Another embodiment of the present invention relates to a computer for producing a three-dimensional model of a molecule or molecular structure, wherein the molecule or molecular structure comprises a three dimensional structure defined by atomic coordinates of a TALL-1 or TALL-1 receptor according to any one the tables of coordinates disclosed
10 herein, or a three-dimensional model of a homologue of the molecule or molecular structure as described above. The computer comprises: (a) a computer-readable medium encoded with the atomic coordinates of the TALL-1 or TALL-1 receptor as described previously herein to create an electronic file; (b) a working memory for storing a graphical display software program for processing the electronic file; (c) a processor coupled to the working memory
15 and to the computer-readable medium which is capable of representing the electronic file as the three dimensional model; and, (d) a display coupled to the processor for visualizing the three dimensional model. The three dimensional structure of the TALL-1 or TALL-1 receptor is displayed or can be displayed on the computer.

Yet another embodiment of the present invention relates to a method to load a
20 therapeutic agent into a carrier for *in vivo* delivery, comprising mixing a therapeutic agent with soluble TALL-1 protein monomers or portions thereof or trimers at a pH below about 7.4 and then raising the pH of the mixture to a pH of about 7.4 or higher to form oligomers of sTALL-1 or portions thereof containing the therapeutic agents for delivery *in vivo*. Also included in the invention are complexes of at least one therapeutic agent and sTALL-1
25 monomers produced by this method. This embodiment relates to the discovery by the present inventors that TALL-1 forms stable oligomers at pH of about 7.4 or higher, but exists as soluble trimers and monomers at lower pH. The therapeutic agent can be any therapeutic agent for which it is desired to use a large carrier such as a TALL-1 oligomer.

Finally, one embodiment of the invention relates to the production of a TNF-family
30 member protein that has been modified by the introduction into the structure of said TNF-family member protein of a structure that is substantially similar to the "flap" structure of a

TALL-1 protein, and the use of such a protein in a method of treatment of a disease or condition that can be regulated by a TNF-family member protein activity.

All publications and patents referenced herein are incorporated herein by reference in their entireties.

5

The following examples are provided for the purpose of illustration and are not intended to limit the scope of the present invention.

Examples

10 Example 1

The following example describes the crystallization of sTALL-1.

Protein expression, purification and crystallization

The cDNA fragment encoding amino acids 134-285 of human TALL-1 (SEQ ID NO:2) was amplified from a TALL-1 full-length cDNA clone (Shu et al., (1999) *J. Leukocyte*
15 *Biology* 65:680-683; Shu et al., (2000) *Proc. Natl. Acad. Sci. USA.* 97:9156-9161) by PCR using a plasmid template and primers containing restriction sites for pET14b (Tagged with His₆)(Novergen). The final clones have been verified by restriction digestion and DNA sequencing. The recombinant plasmid containing the *sTALL-1* gene was transformed to *E. coli* BL21(DE3)pLysS. Enhanced expression of His₆-sTALL-1 was induced by adding IPTG
20 (isopropyl-1-thio-β-D-galactopyranoside) to a 8 L growing culture (37°C) at an OD₆₅₀ of 0.7. After 4 hrs of additional growth, cells were harvested, resuspended in buffer (50mM Tris-HCl, 1mM EDTA, pH 8.0, 300mM NaCl, 5% glycerol and 1mM DTT). After cell lysis through a continuous-flow French press and a low-speed spin, the soluble fraction was loaded onto a Ni²⁺-chelating affinity column and His₆-sTALL-1 was eluted with 100 mM imidazole.
25 His₆-sTALL-1 was loaded onto a MonoQ (Pharmacia) ion-exchange column. After elution with a NaCl gradient, the protein was homogeneous as judged by Coomassie stained SDS-polyacrylamide gels. The His₆ tagged fusion protein was used for final crystallization screen. sTALL-1 (15mg/ml) was crystallized by vapor diffusion against 5 mM β-mercapotethanol, 50 mM Bicine pH 9.0, 150mM NaCl, and 35-38% dioxane. Heavy-atom derivatives were
30 prepared by soaking crystals for 24 hrs in 1mMHgCl₂ and 48 hrs in 1 mM Mersayl, all

dissolved in the crystallization solution. For cryo-crystallography, crystals were soaked in 35% MPD with crystallization buffer for 30 minutes before flash-freezing.

Structure Determination and refinement

Crystals of sTALL-1 diffracted to 3.5Å and have high mosaicity (above 1.0°). Large
5 crystal sizes did not improve the crystal diffraction ability and mosaicity. Altering crystallization conditions did not produce any better results. Finally, mercury treatments (sTALL-1 sample was soaked in 0.1mM HgCl₂ for 10 hours) on the sTALL-1 protein before crystallization were employed. This sample produced crystals at similar conditions as the native protein of sTALL-1. The new crystals diffracted to 2.8Å at an in-house x-ray generator
10 and with reasonable mosaicity (0.5°). The cell dimensions of these crystals are 234Åx234Åx217Å and these crystals are in space group of P6₃22. The molecular weight of the His₆ tagged sTALL-1 is ~20k Da. Assuming all monomers form trimers in solution and crystal packing. It was impossible for the inventors to define the exact solvent contents of the sTALL-1 crystals (Matthews, B.W. (1968) *J. Mol Biol* 33:491-497). Data collection and
15 heavy atom screen were followed. All data sets are processed by DENZO and SCALEPACK (Otwinowski et al., (1997) *Methods Enzymol* 276:307-326). Due to the low sequence homology among TNF ligand family members, it was expected that it might be difficult to find molecular replacement solutions. This turned out to be the case when all available molecular replacement programs were tried. Based on the initial V_m analysis, there are at
20 least 3 trimer or 9 monomers in an asymmetry unit. The inventors tried both monomer and trimer of TNFα, TNFβ, CD40L and TRAIL as initial search models. All of these search models failed to generate a molecular replacement solution. Traditional heavy atom screens were carried out. Two mercury derivatives (Table 1A) were found. Three data sets of sTALL-1 were inputted to the program "SOLVE" (Terwilliger et al., (1987) *Acta*
25 *Crystallogr. A* 43:34-38), and the output map were subjected to solvent flattening in "SOLOMON" (Sheldrick, G. M. (1987) In isomorphous Replacement and Anomalous Scattering, Proc. CCP4 st. W. Wolf, P. R. Evans, A. G. W. Leslie, Eds. (daresbury Laboratory, Warrington, UK) P23). The solvent flattened map was used for model building in program "O" (Jones et al., (1991) *Acta crystallogr. A* 47:110-119). The 3.0Å MIR map
30 of sTALL-1 has excellent quality. All side chains of residues 142-285 were resolved. Tracing and model building were performed with the help of a TNFβ model (Eck et al.,

(1992) *J. Biol. Chem.* **267**:2119–2122). The 10 mercury sites per asymmetric unit and their symmetry related sites lead to two ball-like arrangements in one unit cell ($a=b=234$, $b=217$ Å, $\alpha=90$, $\beta=90$, $\gamma=120$) with each ball containing 60 sites. This feature reminded the inventors of a T=1 virus structure, and also indicates that there are 10 monomers in the asymmetry unit instead of integrate numbers of trimers as was assumed in the calculation of solvent contents initially. The 10 monomers of the asymmetry unit were built independently. sTALL-1 model was first subjected to rigid body refinement and then conventional positional minimization in program “CNS” (Brunger et al., (1998) *Acta. Cryst.* **D54**:905-921) with non-crystallographic symmetry constraints. The output model was then subjected to the “slow cooling” dynamic annealing refinement and a group B-factor refinement. The final refined R factor is 23.6% and R free is 25.2%. It was surprising that there are no mercury atoms in the sTALL-1 structure, which are derived from the pre-HgCl₂ treated sample, although the behavior of the sTALL-1 sample was completely changed after HgCl₂ treatments.

Table 1A Experimental data on crystal structure determination and refinement

	Data Set	Resolution (Å)	R _{merge} (%)	No. of unique reflections		Total Observations		Completeness (%)	Phasing power	No. of Sites	
20	Native	3.0	11.7	66001		334821		95.7			
	HgCl ₂	3.0	14.6	111792		317553		85.2	1.39	10	
	Mersalyl	3.2	11.4	101595		247096		94.5	0.67	10	
	Mean Figure of Merit = 0.37 (for 54518 phased reflections)										
25											
	Refinement Resolution (Å)	20-5.95	4.75	4.15		3.78	3.51	3.30	3.14	3.00	
	No. reflections	7850		7545	7472	7158	6732	6328	4927	3740	51752
	R-factor	23.12		21.06	18.73	22.84	25.08	27.78	31.34	33.67	23.61
	Free R-factor	25.48		21.37	20.45	24.71	26.92	28.07	33.48	35.03	25.22

$R_{\text{merge}} = \sum |I_j - \langle I \rangle| / \sum I_j$ with Bijvoet pairs treated as equivalent for native, as different for derivatives. Total observations, the number of full and partial observations measured with non-negative intensity to the indicates resolution. Completeness, the percentage of possible unique reflections measured with $1/\sigma(I) \geq 0$ to the indicated resolution. Phasing power = $\langle F_H \rangle / E_{\text{rms}}$. No. reflections, the number of reflections used in refinement for each resolution bin. R-factor = $\sum |F_o - F_c| / \sum F_o$ for all amplitudes with $F/\sigma(F) \geq 2$ measured in the indicated resolution bin; the free R-factor is calculated with 5% of the data in each bin.

Overall Structure

The structure of the soluble portion of TALL-1 was determined by the multiple isomorphous replacement method using two mercury derivatives (Table 1A). The electron

density map was improved by solvent flattening. The structure has been refined to an R-factor of 23.6% (R_{free} -factor of 25.2%) against data extending to 3.0 Å resolution in space group $P6_322$, with ten sTALL-1 monomers in the asymmetric unit (unit cell of 234 X 234 X 217 Å). The current model of the sTALL-1 monomer contains residues 142 to 285 (SEQ ID NO:2), with all side-chains well defined (Fig. 1A).

The structure of sTALL-1 consists of two layered antiparallel β strands that form a typical jellyroll-like β sandwich, as with other members of the TNF ligand family (Jones et al., (1989) *Nature* 338:225–228; Eck et al., (1989) *J. Biol. Chem.*, 264:17595-17605; Eck et al., (1992) *J. Biol. Chem.* 267:2119–2122; Karpusas et al., (1995) *Structure* 3:1031–1039; Cha et al., 1999, *Immunity* 11:253-261). Compared to known structures of other family members, the overall structure of sTALL-1 is shorter along the 3-fold axis that generates the trimers (Fig. 1B). This was even more obvious when the trimers of sTALL-1 and TNF α were superimposed, which generated an overall RMSD of 1.9 Å (Fig. 1C). The effect is caused by the shortening of two β strand pairs, CD and EF (Fig. 1A). This is consistent with the fact that the connecting regions that link β strands CD, EF, and GH are the most divergent regions among the TNF family ligands. Two additional unique features are termed "elbow" and "flap" regions (Fig. 1A). The "elbow" region contains a short β hair-pin labeled A' and A". There is a similar region in TRAIL, which is not well defined from the available structures of TRAIL, contrasting with the well ordered β hair-pin of sTALL-1 (Cha et al., 1999, *supra*; Mongkolsapaya et al., 1999, *supra*; Hymowitz et al., 1999, *supra*). The "flap" region is unique to sTALL-1 based on results of sequence alignments and structural comparisons (Fig. 1B, 1D). There is also a disulfide bridge between residue Cys232 on strand E and residue Cys245 on strand F, which is unique for TALL-1, TALL-2, Tweak, and EDA (Bodmer et al., 2000, *supra*).

The interfaces that form the trimer of sTALL-1 mostly consist of layered aromatic residues including Phe194, Tyr196, and Tyr246 from three monomers, which are conserved for all TNF ligand family members. Interestingly, one triple Phenylalanine layer in TNF α , TNF β , CD40L, and TRAIL is replaced by triple Leu282s in sTALL-1 trimer. There are two additional interaction layers of triple residues in sTALL-1 trimer. One consists of residues Gln144 from each monomer, forming a H-bond net. Another layer is formed by three

Leu285 residues from C-terminus of three monomers. The hydrophobic interactions appear to be the main forces driving trimer formation.

The unique "flap" region of sTALL-1 mediates trimer-trimer interactions that lead to a remarkable virus-like assembly of the sTALL-1 trimers. There are 10 sTALL-1 monomers in the asymmetric unit with a space group of $P6_322$ (Fig. 2A). The 10 monomers interact to form virus-like clusters containing 60 sTALL-1 monomers (20 trimers) (Fig. 2B, 2C). Within the unit cell, there are two virus-like clusters. This structure resembles the T=1 virus structures such as satellite tobacco necrosis virus (STNV, PDB ID:2STV) (Jones et al., (1984) *J Mol Biol* 177:735-767). The overall RMSD of main chain is 2.1 Å between sTALL-1 and STNV. In STNV structure, 5 monomers form a pentamer, a virus envelope is built up by 12 pentamers and the interactions among pentamers are mediated by two short helices. Interestingly, when the structure of TNF α was reported, the authors noticed the structural similarity between TNF α and the capsid protein of STNV (Jones et al., (1989) *Nature* 338:225-228). Moreover, they speculated that these two proteins could have evolved from a common ancestor and that TNF α may form a virus-like structure under certain circumstances (Jones et al., (1989) *Nature* 338:225-228). The structure of sTALL-1 is a piece of strong evidence supporting that speculation.

Trimer-trimer interactions

The trimer-trimer interactions are extensive. They not only include hydrogen bond net works and salt bridges, but also hydrophobic contacts. Residues involved in trimer-trimer interactions are not only from the monomer that contributes the "flap" region but also the neighboring monomer as well (Fig. 3A, 3B). Due to the resolution limitation, detailed hydrogen bond networks will not be discussed here. There are three major interaction interfaces that bring two trimers together, two of them are involved in the interactions of two monomers, layer 1 and layer 2 respectively (Fig. 3B). Layer 1 consists of residues Tyr192, Lys252, Glu254, and His218 from one monomer, residues Tyr192', Lys252', Glu254', and His218' from another monomer. The side chains of residue Lys252 and Glu254 form ionic bonds with these of residue Glu254' and Lys252'. These interactions are further strengthened by the hydrogen bond net formed by side chains of all residues from this layer (Fig. 3C, 3D). Interestingly, except residues His218 and His218' from the "flap" regions, all others are from β strands C and F. These interactions could exist in other TNF ligand members that do not

contain a distinguishable "flap" region. Further investigation of the biological consequence of the interaction for other members will be of great interest and importance.

Layer 2 consists of residues Lys216, Glu223, Leu224, Val227, and Leu229 from each monomer (Fig. 3E, 3F). The side chains of residues Lys 216 and Glu223 from one monomer
5 form ionic bonds with Glu223' and Lys216' of another. The side chains of residues Val227, Leu229, part of Lys216 and Glu223' form one hydrophobic core, side chains of Val227', Leu229', part of Lys216' and Glu223 form another. The interaction of residues Leu224 and Leu224' further bolsters the "flap"-"flap" interactions (Fig. 3E, 3F).

The interactions of the third layer are among three monomers, monomer 1, 1' and 2'
10 (Fig. 3G). The side chain of residue Val219 from the "flap" region of monomer 1 interacts with the side chains of residues Ile150 and Leu169 of monomer 2' to form one hydrophobic core. The side chain of residue Phe220 from the "flap" region of monomer 1, side chains of residues Tyr192 and Ile250 from monomer 1', and the side chain of residue Phe172 from monomer 2' form another hydrophobic core. The two hydrophobic cores, which bring three
15 monomers together, are separated by the main chain of the "flap" from monomer 1, (Fig. 3G, 3H). Additional hydrogen bonds formed by three monomers at this region further intensify these tri monomer-monomer interactions. The present inventors believe that these interactions also greatly improve the stability of the traditional trimer, which is formed by monomer 1, 2 and 3 or monomer 1', 2' and 3' (Fig. 3A).

From the previous results discussed in the Background, the trimer is believed to be
20 the functional unit for the TNF ligand family members (Locksley et al., 2001, *supra*; Fesik, 2000, *supra*) (Fig. 1C). Due to the intensive interactions among the sTALL-1 trimers in the crystal structure, the inventors propose that the functional unit comprises more than a single trimer. Based on the crystal structure of sTALL-1, the inventors can isolate several possible
25 sub-clusters that may act as the functional unit *in vivo*. The first is the dimer of trimers (Fig. 3A). The second is the tetramer of trimers (Fig. 5A). The third is the pentamer of trimers, which is formed by encircling trimers (Fig. 5B). Finally, based on the gel-filtration and electron microscopy results (see Example 2), the inventors suggest that the most likely functional unit is the entire virus-like cluster (Fig. 2).

Example 2

The following example demonstrates that the virus-like assembly of TALL-1 exists in solution.

To confirm that the virus-like assembly of sTALL-1 exists in solution and is not the
5 result of a crystal-packing artifact, the present inventors employed two methods, gel-filtration
and electron microscopy. The assembly state of sTALL-1 was investigated on a Superose-6-
gel-filtration column. From the final elution profile, the sTALL-1 sample contains
assemblies with an estimated molecular weight greater than 670,000 Da and smaller than
2,000,000 Da, consistent with the calculated molecular weight of the 60mer virus-like
10 assembly of approximately 1,200,000 Da. To evaluate the stability of the assembly, different
salt concentrations were applied. Three salt concentrations (10mM NaCl, 500mM NaCl, and
1M NaCl in 50mM Tris-HCl buffer at pH 8.0) led to the same sharp elution profile,
consistent with the present inventors' structural information that the trimer-trimer interactions
involve not only electrostatic contacts but also extensive hydrophobic contacts. These results
15 also can be repeated on the Superdex-200 gel-filtration column (Superdex-200 HR10/30,
Pharmacia), although sTALL-1 came out at a void volume (V_0 , 48mL) (data not shown).

These results are not consistent with two published results (Schneider et al., (1999)
J Exp Med. 189:1747-56; Kanakaraj et al., (2001) *Cytokine.* 13:25-31), both of which
claimed that sTALL-1/BAFF existed only as trimers. Comparing these experimental
20 procedures with the published results, the present inventors noticed that the analyses of
sTALL-1/BAFF were carried out at different pHs: pH 6.0 in the Kanakaraj study, pH 7.0 in
the Schneider study, and pH7.4 and above in the present inventors' studies. To evaluate the
influence of pH on the oligomeric state of sTALL-1, gel-filtrations were run using sTALL-1
from bacteria on Superdex-200 at a series of pHs (data not shown). At pH 6.0, sTALL-1
25 exists exclusively as trimers. The ratio of oligomers to trimers at pH 6.5 was 1:2, rose to 1:1
at pH 7.0, and majority are oligomers (30:1) at pH7.2. At pH7.4 the present inventors could
detect only oligomeric sTALL-1 (data not shown). To rule out that sTALL-1 from different
sources may behave differently, purified sTALL-1 from the 293 cell line as well as sTALL-1
produced by a mouse myeloma cell line were analyzed (R&D Systems, Inc., Minneapolis).
30 Both existed exclusively as oligomers at pH7.4 (data not shown). The present inventors
believe that two histidine residues (with $pK_2=6.0$) in the "flap" region may play crucial roles

in the pH dependent association and dissociation property of sTALL-1 (Fig. 3C, 3D). Interestingly, there are trimers as well as monomers of sTALL-1 at Ph7.0.

For electron microscopy, a 5µl of sTALL-1 suspension (5mg/ml) was applied for one minute onto carbon coated grid previously glow discharged. After removing the excess of liquid with a filter paper, the grid was washed with 2 drops of buffer and then negatively
5 stain with saturated uranyl acetate. Electron micrographs were recorded at a magnification of 60,000 x with a Philips CM12 transmission electron microscopy (Philips Electron Optics, Eindhoven, The Netherlands) operating at 100 kV.

The 200 Å diameter virus-like assembly observed in the crystal structure is of
10 sufficient size to be easily detected by electron microscopy. This proved to be the case. The negatively-stained virus-like clusters are clearly observed in electron micrographs after absorption of the sample to a carbon electron microscope (Fig. 4A). The diameter and the shape of the particle observed by electron microscopy is consistent with the crystal structure (Fig. 4B, 4C).

15

Example 3

The following example describes experiments that demonstrate that the "flap" region is involved in the cluster formation by TALL-1.

To further confirm the exact region that leads to the formation of the virus-like cluster
20 and the functional role of the "flap" region, the present inventors constructed a mutated version of sTALL-1 with 8 residues (residue 217 to residue 224 of SEQ ID NO:2) replaced by two glycines at the "flap" region. The truncated sTALL-1 (sTALL-1 217-224) was overexpressed and purified as the native sTALL-1 (see Example 1).

For gel-filtration, wild type sTALL-1 and mutant sTALL-1 (each of 1mg) were
25 loaded onto the Superdex-200 HR 10/60 (Pharmacia) respectively. The wild type sTALL-1 came out at void volume (45-50mL) and the mutant trimers and monomers came out at elution volumes of 78mL and 87mL respectively (data not shown). Direct fresh medium (6 mL) that grow 293 cells was loaded on to the Superdex-200 column. All fractions at different elution volume were concentrated for final western-blot experiment followed the
30 procedure as described (Shu et al., 1999, *supra*), sTALL-1 were detected at elution volume of 48 mL and 80mL respectively (data not shown).

Gel-filtration of the truncated sTALL-1 sample on a Superdex-200 column yielded two peaks with molecular weight of around 60,000 Da and 20,000 Da (data not shown). These peaks consistently appear in the elution profile at all three salt concentrations. These molecular weights correspond to a monomer and a trimer of the truncated sTALL-1. These data demonstrate that after truncation of the "flap" region, the cluster forming property of sTALL-1 is abolished. Furthermore, disruption of the "flap" region also affects trimer formation, which is consistent with the present inventors' observation of a hydrophobic core formed by residues from three monomers at the monomer-monomer interaction interface. Although the "flap" regions of one trimer do not involve in the formation of the trimer itself (Fig. 1C), they do affect the stability of neighboring trimers. These data are also consistent with the observation that for some family members, trimers can be transformed to monomers over a period of time in solution (Cha et al., 1999, *supra*; Corti et al., (1992) *Biochem. J.* 284, 905-910).

From the structural results, the inventors note that the "flap" region is in the corresponding location for the binding of one CRD of the cognate receptors of TNF β and TRAIL. To investigate if the truncation of the "flap" region of sTALL-1 affects its receptor binding capacity and affinity, binding assays of native and truncated sTALL-1 to the cognate receptor BCMA were carried out.

For these experiments, the extracellular domain of BCMA (residues from 1 to 51; SEQ ID NO:6) for BIAcore experiments was overexpressed as GST-BCMA fusion protein on pGEX4T-2 vector in BL-21 strain. Cell preparation and protein purification procedure are similar to that of sTALL-1. Briefly, harvest cells were broken through a French press and low speed spin. The soluble fraction was loaded onto a GST affinity beads. After intensive wash with binding buffer, thrombin was added for 24 hrs incubation. Supernatant that contained the extracellular domain of BCMA was loaded onto a MonoQ column and eluted with NaCl gradient. The protein is above 99% pure after MonoQ.

First, BCMA-transfected 293 cells were incubated with control medium, wild type sTALL-1 in the control medium, and mutant sTALL-1 in the control medium. Cell staining was performed by sequential incubation (each 40 min) with anti-His₆ mAb and RPE-conjugated goat anti-rabbit IgG in staining buffer. The fluorescence was measured by using a Becton Dickinson FACScan flow cytometer (Shu et al., 2000, *supra*) (data not shown).

Flow cytometry analysis results showed that the truncated sTALL-1 and the wild sTALL-1 had similar binding capacity to cells that overexpressed entire BCMA (data not shown).

The binding affinity of BCMA for the wild type versus mutant sTALL-1 was further determined using BIAcore surface plasmon resonance. To obtain the true affinity of the receptor without the confounding effect of multivalent binding, the polyvalent TALL-1 proteins were immobilized in the instrument flow cells and the soluble monomeric BCMA was injected in the mobile phase. The KD of the interaction was calculated by Scatchard analysis of the equilibrium data (data not shown). Briefly, wild type multimeric sTALL-1 and mutant trimeric sTALL-1 were immobilized (11,000 and 3000 RU, respectively) in separate flow cells of a CM-5 BIAcore Biosensor chip using standard amine coupling reagents. Various concentrations (10-1000nM) of monomeric BMCA were injected for one min at a flow rate of 20 μ l/min in a buffer of 100mMNaCl, 20mM Tris-HCl pH8.0 and 0.005% P20 detergent and the binding kinetics recording. For base line calculations the same BMCA samples were injected in a control flow cell with no protein immobilized. Receptor binding kinetics were too rapid for accurate measurements of on rates and off rates, but equilibrium binding values (Rmax) were used to calculate the overall affinity of receptor binding by Scatchard analysis.

Results showed that the receptor bound to the two ligands with very similar affinity, indicating the removal of the TALL-1 flap did not alter its affinity for its receptor. An affinity of about 100nM for this monomer interaction is consistent with reports of 0.1-1.0nM apparent affinities for trivalent interactions of TNF family members with immobilized or cell bound receptors. Therefore, it is clear that the "flap" region is not involved in receptor binding.

To understand the functional role of the "flap" region, transfection assays of both wild type and truncated sTALL-1 were performed. Luciferase reporter gene assays were carried out as described (Shu et al., 2000, *supra*). Briefly, 293 cells were transfected with 0.5 μ g of NF- κ B luciferase reporter plasmid and increased amounts of an expression plasmid for BCMA. Fourteen hours after transfection, cells were treated wild type sTALL-1, mutant sTALL-1, or left untreated for 7 hrs and luciferase reporter assays were performed.

Wild type sTALL-1 gave a dose dependent activation of NF- κ B in reporter gene assays. Truncated sTALL-1 was defective in activating NF- κ B at a variety of BCMA

concentrations (data not shown). These data demonstrated that, despite the normal binding of the truncated sTALL-1 to the receptor, the "flap" region is essential for the proper function of sTALL-1 *in vivo*.

To avoid possible artificial results brought in from the 293 cell transfection assay system, B lymphocyte (from human peripheral blood of health donors) proliferation stimulation by wild type sTALL-1 and mutant sTALL-1 were carried out. B cell proliferation assays followed the procedure as described (Shu et al., 2000, *supra*). Briefly, human peripheral B lymphocytes were purified from peripheral blood of health donors. Purified B lymphocytes were seeded on 96-well dishes and treated with indicated reagents for 40h. Cells were pulsed for an additional 10h with [³H]thymidine. Incorporation of [³H]thymidine was measured by liquid scintillation counting (data not shown). These costimulation assays indicated that wild type sTALL-1, but not mutant sTALL-1 significantly ($P < 0.01$) stimulate B lymphocyte proliferation (data not shown).

Example 4

The following example demonstrates that sTALL-1 clusters exist under physiological conditions.

To assess whether the sTALL-1 cluster exists *in vivo*, medium from sTALL-1 overexpressing 293 cells was collected and loaded onto a gel-filtration column (Superdex-200 HR 10/60, Pharmacia) eluting with PBS buffer at pH7.4. The eluted fractions were then subjected to western-blot analysis. sTALL-1 exists both as clusters and trimers, judging from the elution volumes (data not shown), indicating that sTALL-1 clusters could exist under physiological condition. To find out if there is an equilibrium between clusters and trimers, the two peaks corresponding to clusters and trimers were collected and concentrated and finally applied back to the same gel-filtration column. For the cluster peak fraction, the same elution profile was obtained, a single sharp and symmetric peak at the cluster position, there was no detectable sTALL-1 at the trimer position. On the other hand, the trimer peak fraction generated two peaks corresponding to the cluster and trimer positions (data not shown). It is clear from these results that sTALL-1 predominantly in the cluster state rather than a trimer in solution at pH7.4. The process of cluster formation from trimers is irreversible.

Current data indicate that TNF family ligands function as trimers binding to the cognate receptors. The recruitment of receptors leads to clustering of the cytoplasmic domains, which in turn stimulates the recruitment of adaptor proteins and other downstream partners. The virus-like cluster of sTALL-1, which the present inventors have shown is
5 required for function, contradicts this paradigm. There is one possible mechanism that may involve the activation of the downstream pathway. The virus-like cluster of sTALL-1 could recruit numerous receptor trimers. This increase of local concentration of the receptors could facilitate signaling through the cellular membrane to the cytoplasm of the cell. The clustering of numerous cytoplasmic domains could lead to signal amplification by recruiting
10 downstream elements. This model resembles the well-characterized SMAC (SuperMolecular Activation Cluster) complexes (Monks et al., (1998) *Nature* 395:82-6), in which multiple copies of T cell receptors, peptide bound major histocompatibility complex molecules, other related accessory proteins, and their counter-receptors from T cells and antigen presenting cells gather together to form clusters that resemble neural synapses (Monks et al., (1998)
15 *Nature* 395:82-6).

Example 5

The following example describes an analysis of other TNF family members for the "flap" region discovered in TALL-1.

20 The present inventors' structural and functional analysis indicates that the sTALL-1 "flap" region mediates the cluster assembly formation, and is essential for the activation of NF- κ B. To determine whether this flap region exists in other family members, the inventors performed a structure-based sequence alignment of all available 18 family members (TALL-1, TRAIL, TNF β , CD40L, TNF α , RANKL, APRIL, FasL, LT β , CD30L, CD27L, OX40L,
25 4-1BBL, EDA-A1, EDA-A2, AITRL, VEGI, LIGHT, TWEAK). The four known TNF ligand structures (Jones, et al., 1989; Eck, et al., 1989; Eck, et al., 1992; Karpusas, et al., 1995; Cha, et al; 1999) were superimposed on sTALL-1 by the program Dali (Holm and Sander, 1993). To search possible "flap" regions in other members, the sequences that span β strands D and E from the five structures were aligned based on the structural
30 superimpositions. The hydrophobic pattern was obvious in this region after alignments (data not shown). For example, residues Met208 and Ile212 on strand D must be hydrophobic for

the proper formation of the hydrophobic core. A similar situation exists for residues Leu229 and Phe230 on strand E. All other family members with unknown structures were aligned according to this pattern (data not shown).

From the alignment results, it is obvious that the loop region connecting β strands D and E is diverse. sTALL-1 has the longest sequence in this region. It is not clear whether any other member will form the "flap" region as in sTALL-1. Further structural, functional, and analysis of solution properties for each individual family member will be necessary. The fact that other family members bind receptors with three or more CRDs, the possibility of virus-like clusters as the functional unit may be limited. This does not exclude the possibility of weak trimer-trimer interaction for functional purpose. If the "flap" exists for functional purposes in other family members, it is possible that only CRD1 of their cognate receptors takes part in the recognition. One interesting observation is that there is a complementary region in receptors of TNF β and TRAIL that interacts with this region in a fashion similar to the trimer-trimer interactions of sTALL-1. Therefore, it is possible that this region may mediate intermolecular interactions *in vivo*, and the corresponding receptor may compete for binding to this region.

Example 6

The following example demonstrates the structure determination of sTALL-1 complexed with eBCMA and eBAFF-R.

Protein expression, purification and crystallization

Protein expression, purification, and crystallization for sTALL-1 is as described in Example 1. BAFF-R/BR3 is cloned according to the published reports (Thompson et al., 2001, *supra*; Yan et al., 2001, *supra*). The extracellular domains of BCMA and BAFF-R used for the experiments were overexpressed as GST fusion protein on pGEX4T-2 vector in the BL-21 strain (Example 1). Cell preparation and protein purification procedure are similar to that of sTALL-1. Briefly, harvest cells were broken through a French press and low speed spin. The soluble fraction was loaded onto a GST affinity beads. After intensive wash with binding buffer, thrombin was added for 24 hrs incubation. The supernatants containing the extracellular domain of BCMA or BAFF-R were loaded onto a MonoQ column and eluted with NaCl gradient. The protein is above 99% pure after MonoQ.

Structure Determination and refinement

For complex crystal preparations, sTALL-1 crystals were harvested after two weeks. sTALL-1 crystals were transferred to a stable soaking solution, which contains 40% dioxane, 1mM correspondent receptors with 100mM Bicine pH 9.0. After overnight soaking, crystals
5 were transferred to the same cryo-protectant buffer system for sTALL-1 crystals (Example 1).

Data sets for both complexes were first collected on the house x-ray generator. Crystals of complexes both diffract to 3.0Å. A 2.6Å data set of eBCMA and sTALL-1 complex were collected at APS. All data were processed by DENZO package (Otwinowski
10 et al., 1997, *Methods Enzymol* 276:307-326). Structures of the complexes were solved by fourier difference using the sTALL-1 model (Example 1). After one run of minimization of sTALL-1 model in CNS (Brunger et al., 1998, *supra*) with new data sets, 2Fo-Fc and Of-Fc maps were calculated. Models were built in O (Jones et al., 1991, *Acta Cryst.* A47:110-119), and finally refined in CNS.

Discussion of Results

15

Crystals of the soluble portion of TALL-1 (sTALL-1) with extracellular domains of BCMA (eBCMA) and BAFF-R (eBAFF-R) were obtained by diffusing the receptor fragments into the sTALL-1 crystals (see above). The structures of both complexes were determined by Difference Fourier using the available sTALL-1 model (Figs 6A-6C and Table
20 1B). The structure of sTALL-1 with eBCMA has been refined to an R-factor of 20.9% (R_{free} -factor of 23.4%) against data extending to 2.6 Å resolution in space group $P6_322$, with ten sTALL-1 monomers and seven entire and one partial eBCMA molecules in the asymmetric unit (unit cell of 234 X 234 X 217Å) (Table 1B). Due to crystal packing, another two receptor binding sites were left unoccupied. Similar results are true for the sTALL-1 and
25 eBAFF-R complex with a final resolution of 3.0 Å (Table 1B). The current model of the eBCMA monomer contains residues 5 to 43 (Fig. 6B). The model of the eBAFF-R monomer contains residues 16 to 58 (Fig. 6C). All figures are prepared by RIBBON (Carson, M. Ribbon models of macromolecules. *J. Mol. Graphics* 5, 103-106 (1987).

Table 1B Experimental data on crystal structure determination and refinement

	Data Set	Resolution (Å)	R_{merge} (%)	No. of unique reflections	Total Observations	Completeness (%)	R-factor	Free R-factor
5	BCMA	2.6	12.9	97672	1056950	94.0	20.9	23.5
	BAFF-R	3.0	13.8	63376	318144	93.3	24.5	26.1

10 $R_{\text{merge}} = \sum |I_i - \langle I \rangle| / \sum I_i$ with Bijvoet pairs treated as equivalent. Total observations, the number of full and partial observations measured with non-negative intensity to the indicates resolution. Completeness, the percentage of possible unique reflections measured with $I/\sigma(I) \geq 0$ to the indicated resolution R -factor = $\sum |F_o - F_c| / \sum F_o$ for all amplitudes with $F/\sigma(F) \geq 0$ measured; the free R-factor is calculated with 5% of the data.

Overall structure

15 The space group of the TALL-1 crystals remained $P6_322$ with the same cell dimensions with or without bindings of the receptors. There are two virus-like clusters in one unit cell. Each cluster has 60 copies of sTALL-1, 42 fully occupied eBCMA or eBAFF-R, 6 partial copies of eBCMA or eBAFF-R. There are 12 copies of sTALL-1 free of receptors due to crystal packing. All receptors are located on the outer-extreme shell, which
20 expands the ball-like shell another ~ 20 Å in each direction. The overall arrangement of the receptors on the shell looks like a sunflower with receptors as flower petals and sTALL-1 as a seed bed (Fig. 7A-7D). Molecules marked red are missing in the complex structure. Molecules marked blue are partially occupied. The conformational change in sTALL-1 is negligible before or after receptor binding, which is the only similarity between this
25 interaction and that of other TNF family members.

Structure of eBCMA

As predicted, eBCMA contains two modules, one is A1 module consisting of three beta strands with strand 1 and strand 3 linked by the only disulfide bridge (Naismith et al., 1998, *supra*). The other module is C2-like (the two disulfide bridges formed as CysI-CysIV
30 and CysII-CysIII), but the disulfide arrangement is the same as a typical of B2 module (the two disulfide bridges formed as CysI-CysIII and CysII-CysIV) (Naismith et al., 1998, *supra*) (Fig. 6B). For clarity, the present inventors temporarily termed it D2 for its difference from C2 and B2. There are two short helices in the D2 module that are located just at the N-terminus and the C-terminus the module, one is from CysI to CysII and another one is from
35 CysIII to CysIV. The latter helix extends further after the disulfide bridge and forms a 4 turns-long helix, which is unique when compared to all known TNF receptor structures. The

arrangement of A1 and D2 of eBCMA is similar to that of A1 and C2 in TNF-R1 (Naismith et al., 1998, *supra*), A1 and D2 form a saddle-like architecture with each module as half of the saddle and the unique helix as the “rider” (Fig. 6B). From the initial 2Fo-Fc and Fo-Fc maps, all seven copies of A1 modules are very rigid with temperature factors similar to that of sTALL-1, and most side-chains are ordered. Interestingly, the partial copy of the eighth eBCMA only contains the A1 module. The RMSD is 0.2Å of eight A1 modules in the asymmetry unit. In contrast, the D2 modules are relatively flexible, especially the region between CysII and CysIII. The RMSD is 1.5Å of the seven D2 modules in the asymmetry unit. The eighth D2 is almost completely disordered except for the region between CysI to CysII.

Structure of eBAFF-R

To the present inventors' surprise, eBAFF-R has similar fold as eBCMA, although it is predicted that eBAFF-R contains only one C2 or X2 module (Thompson et al., 2001, *supra*; Yan et al., 2001, *supra*; Bodmer et al., 2002, *supra*) (Fig. 6C). The structure of eBAFF-R shows that it also contains two modules, A1 and the C2-like modules. The A1 module contains three cysteines, only two of them (Cys19 and Cys32) form the typical disulfide bridge. Interestingly, the C2-like module in eBAFF-R only contains one cysteine and it is impossible for it to form disulfide bridges. From the initial 2Fo-fc and Fo-Fc maps, the inventors did find some connecting density at the equivalent CysI-CysIII and CysII-CysIV disulfide bridge positions in the D2 module of eBCMA. Actually, this density represents a H-bond formed between the only Cys35 (location CysI) and the side chain of Ser49 at the equivalent location as the CysIII in eBCMA. Similarly, the Arg39 at the equivalent location CysII forms a H-bond with the oxygen of the main chain of Ala52 at the equivalent location CysIV in eBCMA. These two hydrogen bonds replace the two disulfide bridges in the D2 module of eBCMA, so we termed this module in eBAFF-R as D0. Although the distance between the free Cys24 in A1 and the one (Cys35) in D0 are ideal for disulfide bridge formation (6.5Å), the side chain orientation of the former cysteine is away from the latter cysteine due to the main chain constraint. Except for what is discussed above, all other structure features are highly similar as described for eBCMA in previous section.

Comparison of eBCMA with eBAFF-R

The sequence homology between eBCMA and eTACI (extracellular domain of TACI) is obvious. This is not true for eBCMA and eBAFF-R or between eTACI and eBAFF-R. The structures of eBCMA and eBAFF-R allowed us to perform a structural based sequence alignment of eBCMA, eBAFF, and eTACI. The inventors found that a strong pattern of similarity emerges (Figs. 8A-8B). As mentioned above, eBCMA and eBAFF-R have a similar saddle-like fold, the RMSD between two equivalent and best defined structures from each group is 1.9Å. (Fig. 8A). A1 modules from eBCMA and eBAFF-R (with RMSD 0.5Å) are almost identical judged by both primary sequence alignment and structure superposition (Figs. 8A-8B). The D2 from BCMA and D0 from BAFF-R are also very similar structurally, although the sequence similarity is poor.

The high sequence similarities between the two CRDs in TACI and CRDs in BCMA or BAFF-R lead the present inventors to predict that each CRD of TACI contains one A1 module and one D2 module. Interestingly, another TNF receptor member Fn14, which also just contains one CRD, was predicted to contain one A1 module and one C2 module (Bodmer et al., 2002, *supra*). From the sequence alignment result presented herein, it could contain either D2 or C2 (Fig. 8B). In any event, without being bound by theory, the present inventors propose that the interaction mode between Fn14 and its ligand TWEAK are similar to what they found in the complexes of eBCMA and eBAFF-R with sTALL-1 (see below).

Interactions of sTALL-1 and eBCMA

The interactions between sTALL-1 and eBCMA are mostly in a one to one mode, that is one monomer of the receptor to one monomer of the ligand. The slightly tilted saddle-like receptor is sitting on a horseback-like groove, which is formed by four loops from the ligand, two (connection regions for strands GH and A'A) at one side and two (connection regions for strands CD and EF) at the other (Figs. 9A-9C). This mode of interaction is dramatically different from that seen within other TNF family members, in which one elongated receptor binds to the cleft formed by two ligands. The interactions of eBCMA with sTALL-1 include hydrogen bonds, salt bridges, and, most importantly, hydrophobic contacts. There are 21 total residues involved, 9 from eBCMA (Tyr13, Asp15, Leu17, Leu18, His19, Ile22, Leu26, Arg27, and Pro34), 8 from the primary ligand (Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266), and 4 from the second ligand (Leu200, Leu240, Asp273, Asp275) of the trimer (Fig. 9D). The overall interactions can be divided into four groups.

First, Leu17, Leu18 from eBCMA, and Tyr163, Leu211, Ile233, Pro264 from the primary ligand, and Leu200 from the second ligand form the first hydrophobic core. Interestingly, Tyr163, Leu200, Ile233, and Pro264 are located on a curved track, the joining of Leu17 from eBCMA makes a perfect hydrophobic curved track, just like a key and lock pair. Leu211 from the primary ligand and Leu18 from eBCMA further strengthen the contacts (Fig. 9E).

Second, Ile22, Leu26 from eBCMA, Tyr206 from the primary ligand, and Leu240 from the second ligand form the second hydrophobic core (Fig. 9F).

Third, Asp15 from eBCMA and Arg265 from the primary ligand forms one salt bridge, and Arg27 from eBCMA and Glu266 from the primary ligand forms another. It is also possible that there is a hydrogen bond between Tyr206 from the primary ligand and Tyr13 from the eBCMA (Fig. 9G).

Fourth, His19 from eBCMA, forms a water mediated interactions with Asp 273, Asp275 from the second ligand.

15 *Interactions of sTALL-1 and eBAFF-R*

The interactions between sTALL-1 and eBAFF-R are similar to those between sTALL-1 and eBCMA, although details are slightly different. The interactions also include hydrogen bonds, salt bridges, and hydrophobic contacts. There are totally 22 residues involved, 9 from eBAFF-R (Asp26, Leu28, Val29, Arg30, Val33, Leu37, Leu38, and Arg42, and Pro45), 8 from the primary ligand (Tyr163, Tyr206, Leu211, Arg 231, Ile233, Glu238, Pro264, Arg265), 4 from the second ligand (Leu200, Leu240, Asp273, Asp275), and one from a flap region of the neighboring trimer (Asp222) (Fig. 9H). The overall interactions also can be divided into four groups.

First, Leu28, Val29 from eBAFF-R, Tyr163, Leu211, Ile233, Pro264 from the primary ligand, and Leu200 from the second ligand form the first hydrophobic core. Compared to eBCMA, Val29 in eBAFF-R replace the equivalent Leu18. The side-chain is shortened, and this could reduce the strength of the contact.

Second, Val33, Leu37, Leu38 and Pro45 from eBAFF-R, Tyr206 from the primary ligand, and Leu240 from the second ligand form the second hydrophobic core (Fig. 9I). Val33 in eBCMA is Ile22. Leu38 is additional for eBAFF-R.

Third, Asp26 from eBCMA and Arg265 from the primary ligand form a salt bridge. The replacement of Arg27 in eBCMA with Leu38 in eBAFF-R eliminates a salt bridge with Glu266 from the primary ligand. However, there is one new salt bridge between Arg42 from eBAFF-R and Glu238 from the primary ligand.

5 Fourth, Arg30 from eBAFF-R, Arg231 from the primary ligand, Asp 273, Asp275 from the second ligand, and Asp222 from the third ligand (from the neighboring trimer) form a complicated salt bridge network (Fig. 9J). The long side chain of Arg30 from eBAFF-R (His19 in eBCMA) is in a position to make contacts with either Asp275 or Asp222. The well-defined electron density of the Arg30 side chain from eBAFF-R in the initial difference
10 maps suggests that these are strong interactions, that might considerably strengthen the eBAFF-R and sTALL-1 binding.

Structural basis of BAFF-R discriminating between APRIL and TALL-1

Two publications that initially reported the cloning of BAFF-R/BR3 receptor claimed that BAFF-R specifically binds to TALL-1 but not APRIL/TALL-2 (Thompson et al., 2001,
15 *supra*; Yan et al., 2001, *supra*). Furthermore, APRIL has very low abundance in all tissues, and was proposed to be dispensable for B-cell maturation (Hahne et al., 1998, *J. Exp. Med.* 188:1185-1190; Schneider et al., 2001, *J. Exp. Med.* 194:1691-1697). It was also predicted that there may be an additional and more specific receptor for APRIL (Ware et al., 2000, *J. Exp. Med.* 192:F35-37. The binding affinity of APRIL to BCMA and TACI are similar to
20 that of TALL-1 to BCMA and TACI (Yu et al., 2000, *Nature immunol.* 1:252-256). From the above structural analysis, eBCMA and eBAFF-R have nearly identical three-dimensional structure. Furthermore, the interactions between eBCMA and sTALL-1 are also highly conserved in the interactions between eBAFF-R and sTALL-1. Without being bound by theory, the present inventors propose that these interactions are also conserved for TACI and
25 sTALL-1. Given these structural similarities, one can investigate begin to investigate how BAFF-R can discriminate between TALL-1 and APRIL.

APRIL was modeled based on the sTALL-1 structure, benefiting from the high primary sequence homology between TALL-1 and APRIL (Shu et al., 1999, *supra*). The most obvious difference between sTALL-1 and APRIL is in the "flap" region (6 residues) of
30 sTALL-1, which is missing in APRIL (Shu et al., 1999, *supra*). In Example 3, the inventors described a mutated version of sTALL-1 with 8 residues replaced by two glycine residues,

which was not functional in transfection assays or in the B-cell stimulation assays, but had a binding affinity to its receptors similar to that of the native sTALL-1. The structure of this mutated sTALL-1 has been determined at 1.7 Å resolution by MIR method by the present inventors, and is almost identical to the sTALL-1 except missing the flap (Liu and Zhang,
5 unpublished). This mutated sTALL-1 is a close model of APRIL.

The final built model of APRIL was imported to the minimization program in CNS (Brunger et al., 1998, *Acta. Cryst.* D54:905-921). The output coordinates were superimposed on the sTALL-1 structure (Fig. 10A). Detailed interactions of eBAFF-R and APRIL were briefly analyzed. All residues from eBAFF-R that are involved in the interactions between
10 the eBAFF-R and sTALL-1 are displayed (Fig. 10B). All equivalent residues in APRIL, which are close to the receptor binding surface in sTALL-1 are also showed (Fig. 10B). To the present inventors' surprise, the interactions are extremely similar to those found in the complexes of eBCMA or eBAFF-R with sTALL-1. First, the first hydrophobic core that was described in the two previous complexes still exists, including residues, Leu28, Val29 from
15 the eBAFF-R and residues, Val133, Thr177, Val181, Ile197, Pro230 from the primary APRIL molecule, and residues Leu170, Tyr208 from the secondary APRIL. The Tyr208 is additional for APRIL, which may strengthen the hydrophobic contact (Fig.10B). Second, the second hydrophobic core consists of residues Val33, Leu37, Leu38, Pro45 from the eBAFF-R, residue Phe176 from the primary ligand, and residue Arg206 from the secondary ligand.
20 Compared to the interactions between eBAFF-R and sTALL-1, Tyr206 from the primary ligand is changed to Phe176. Leu240 from the second ligand is changed to Arg206. The former change could also strengthen the hydrophobic contacts (Fig. 10B). Third, the major salt bridge formed by residue Asp26 from the receptor and residue Arg231 from APRIL is conserved (not shown). These comparisons have lead the present inventors to conclude,
25 without being bound by theory, that APRIL could be able, at least under some conditions, to bind to BAFF-R, which is contradictory to previous reports (Thompson et al., 2001, *supra*; Yan et al., 2001, *supra*).

To test these predictions, the binding affinity of eBAFF-R for the sTALL-1 versus APRIL was determined using BIACore surface plasmon resonance. To obtain the true
30 affinity of the receptor without the compounding effect of multivalent binding, the polyvalent sTALL-1 proteins and APRIL were immobilized in the instrument flow cells and soluble

monomeric eBAFF-R and eBCMA were injected in the mobile phase. The k_{ds} of the interaction were calculated from the kinetic binding data (data not shown). To avoid nonspecific binding, the inventors analyzed a series of binding curves at pH7.5, pH8.0, pH8.5, and pH9.0. To the inventors' surprise, it was discovered that eBAFF-R only binds APRIL at pH8.5 or higher, and the binding affinity increases coupled with pH increasing (data not shown). To address the new property of eBAFF-R binding to APRIL, a serial truncation, a point mutation, and a combination of the two were introduced (data not shown). All protein versions with these mutations were subjected to BIAcore binding assays. The inventors only found two versions of eBAFF-R mutations to have significant binding affinity to APRIL at pH7.5; they are (1) residues 12-62 (SEQ ID NO:8) (substitutions at R30H, H31A), and (2) residues 12-62 (SEQ ID NO:8) (substitutions at V29L, V33I) respectively. The rationale for the first version is that the highly positive charged N-terminal of eBAFF-R and Arg30 prevent eBAFF-R from binding to APRIL, which also has a high positive charge on its surface, based on the present inventors' model and others, with a predicted PI of 9.4. Interestingly, the homologues with only a 1 to 12 truncation or only a R30H, H31A double mutation, do not change the binding property. For the second version, which partially mimics eBCMA (although Arg30-His31 is still in the version), the inventors believe, without being bound by theory, that strong hydrophobic interaction forces introduced by the long side-chain of Leu and Ile overcome the inhibitory role of Arg30-His31 (data not shown). Interestingly, eBCMA loses binding affinity to APRIL at pH6.0 (data not shown). These data further suggested that positively charged His19 (equivalent to position Arg30 of eBAFF-R) also plays an inhibitory role for the binding of eBCMA to APRIL.

The inventors have speculated that APRIL could play a decoy ligand role. These current data show that eBAFF-R does not bind to APRIL at normal physiological conditions (at pH7.5), and so the mechanism of APRIL competitively binding with TALL-1 for BAFF-R binding is not expected to be true under normal physiological conditions. From the modeling results, the inventors found that residues that are involved in trimerization are absolutely conserved (data not shown). This information suggested if TALL-1 and APRIL are expressed in the same environment and at the same time, they could form heterotrimers. It has turned out to be true that patients with autoimmune diseases have detectable heterotrimers of TALL-1 and APRIL. As the inventors have concluded previously herein,

sTALL-1 trimers alone (e.g., absent clustering of trimers) can not trigger the signal transduction of its cognate receptors. Heterotrimers of TALL- 1 and APRIL could not form the virus-like cluster as sTALL-1 alone due to the lack of the "flap" region in APRIL. Considering the fact that overexpression of TALL-1 could lead to the abnormal stimulation
5 of B-cell and finally the development of autoimmune disease, APRIL may be serving as a balancer, reducing the opportunity for sTALL-1 to form the active cluster. This role is similar to the decoy death receptors, which are essential for cells to survive ((Cha et al., 1999, *Immunity* 11:253-261; Mongkolsapaya et al., 1999, *Nat. Struct. Biol.* 6:1048-1053; Hymowitz et al., 1999, *Mol. Cell* 4:563-571). Finally, knock-out data for APRIL showed that
10 mice die at early embryonic stages, which indirectly suggests an important role of APRIL (Mackay et al., 2002, *TRENDS in Immunology* 23:113-115).

In this example, two novel structural modules of TNF receptors (D2 and D0) are revealed from the structures of sTALL-1 complexed with eBCMA and with eBAFF-R. The interaction modes of the eBCMA and eBAFF-R with sTALL-1 are completely different from
15 those found in the other TNF family members, containing at least two CRDs that bind to the cleft regions formed by two ligands. For the interactions described here, one saddle-like receptor mostly makes a one to one interaction with its ligand at the extreme end of the ligand. The difference exists not only in the CRD structure but also in the binding locations and modes. The structural based sequence alignment indicates that similar interaction modes
20 may also exist in the interaction between Fn14 to TWEAK, another TNF family couple. Furthermore, modeling analysis contributes to the proposal that APRIL could be a critical decoy ligand functioning in a similar way to the decoy death receptors.

TABLE 2

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REMARK 3
REMARK 3 REFINEMENT.
5  REMARK 3 PROGRAM : CNS 0.9
REMARK 3 AUTHORS : BRUNGER, ADAMS, CLORE, DELANO,
REMARK 3 GROS, GROSSE-KUNSTLEVE, JIANG,
REMARK 3 KUSZEWSKI, NILGES, PANNU, READ,
REMARK 3 RICE, SIMONSON, WARREN
10  REMARK 3
REMARK 3 DATA USED IN REFINEMENT.
REMARK 3 RESOLUTION RANGE HIGH (ANGSTROMS) : 3.00
REMARK 3 RESOLUTION RANGE LOW (ANGSTROMS) : 19.99
REMARK 3 DATA CUTOFF (SIGMA(F)) : 2.0
15  REMARK 3 DATA CUTOFF HIGH (ABS(F)) : 172269.28
REMARK 3 DATA CUTOFF LOW (ABS(F)) : 0.000000
REMARK 3 COMPLETENESS (WORKING+TEST) (%) : 79.3
REMARK 3 NUMBER OF REFLECTIONS : 54509
REMARK 3
20  REMARK 3 FIT TO DATA USED IN REFINEMENT.
REMARK 3 CROSS-VALIDATION METHOD : THROUGHOUT
REMARK 3 FREE R VALUE TEST SET SELECTION : RANDOM
REMARK 3 R VALUE (WORKING SET) : 0.236
REMARK 3 FREE R VALUE : 0.252
25  REMARK 3 FREE R VALUE TEST SET SIZE (%) : 5.1
REMARK 3 FREE R VALUE TEST SET COUNT : 2771
REMARK 3 ESTIMATED ERROR OF FREE R VALUE : 0.005
REMARK 3
REMARK 3 FIT IN THE HIGHEST RESOLUTION BIN.
30  REMARK 3 TOTAL NUMBER OF BINS USED : 6
REMARK 3 BIN RESOLUTION RANGE HIGH (A) : 3.00
REMARK 3 BIN RESOLUTION RANGE LOW (A) : 3.19
REMARK 3 BIN COMPLETENESS (WORKING+TEST) (%) : 48.7
REMARK 3 REFLECTIONS IN BIN (WORKING SET) : 5219
35  REMARK 3 BIN R VALUE (WORKING SET) : 0.331
REMARK 3 BIN FREE R VALUE : 0.344
REMARK 3 BIN FREE R VALUE TEST SET SIZE (%) : 5.0
REMARK 3 BIN FREE R VALUE TEST SET COUNT : 275
REMARK 3 ESTIMATED ERROR OF BIN FREE R VALUE : 0.021
40  REMARK 3
REMARK 3 NUMBER OF NON-HYDROGEN ATOMS USED IN REFINEMENT.
REMARK 3 PROTEIN ATOMS : 0
REMARK 3 NUCLEIC ACID ATOMS : 0
REMARK 3 HETEROGEN ATOMS : 0
45  REMARK 3 SOLVENT ATOMS : 0
REMARK 3
REMARK 3 B VALUES.
REMARK 3 FROM WILSON PLOT (A**2) : $BW
REMARK 3 MEAN B VALUE (OVERALL, A**2) : 30.6
50  REMARK 3 OVERALL ANISOTROPIC B VALUE.
REMARK 3 B11 (A**2) : 1.50
REMARK 3 B22 (A**2) : 1.50
REMARK 3 B33 (A**2) : -3.00
REMARK 3 B12 (A**2) : 5.87
55  REMARK 3 B13 (A**2) : 0.00
REMARK 3 B23 (A**2) : 0.00
REMARK 3
REMARK 3 BULK SOLVENT MODELING.
REMARK 3 METHOD USED : FLAT MODEL
60  REMARK 3 KSOL : 0.285354
REMARK 3 BSOL : 10 (A**2)
REMARK 3
REMARK 3 ESTIMATED COORDINATE ERROR.
REMARK 3 ESD FROM LUZZATI PLOT (A) : 0.40
65  REMARK 3 ESD FROM SIGMAA (A) : 0.62
REMARK 3 LOW RESOLUTION CUTOFF (A) : 5.00

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REMARK 3
REMARK 3 CROSS-VALIDATED ESTIMATED COORDINATE ERROR.
REMARK 3 ESD FROM C-V LUZZATI PLOT (A) : 0.42
REMARK 3 ESD FROM C-V SIGMAA (A) : 0.65
5 REMARK 3
REMARK 3 RMS DEVIATIONS FROM IDEAL VALUES.
REMARK 3 BOND LENGTHS (A) : 0.009
REMARK 3 BOND ANGLES (DEGREES) : 1.6
REMARK 3 DIHEDRAL ANGLES (DEGREES) : 26.8
10 REMARK 3 IMPROPER ANGLES (DEGREES) : 0.84
REMARK 3
REMARK 3 ISOTROPIC THERMAL MODEL : RESTRAINED
REMARK 3
REMARK 3 ISOTROPIC THERMAL FACTOR RESTRAINTS. RMS SIGMA
15 REMARK 3 MAIN-CHAIN BOND (A**2) : NULL ; NULL
REMARK 3 MAIN-CHAIN ANGLE (A**2) : NULL ; NULL
REMARK 3 SIDE-CHAIN BOND (A**2) : NULL ; NULL
REMARK 3 SIDE-CHAIN ANGLE (A**2) : NULL ; NULL
REMARK 3
20 REMARK 3 NCS MODEL : CONSTR
REMARK 3
REMARK 3 NCS RESTRAINTS. RMS SIGMA/WEIGHT
REMARK 3 GROUP 1 POSITIONAL (A) : NULL ; NULL
REMARK 3 GROUP 1 B-FACTOR (A**2) : NULL ; NULL
25 REMARK 3
REMARK 3 PARAMETER FILE 1 : CNS_TOPPAR/protein_rep.param
REMARK 3 PARAMETER FILE 2 : CNS_TOPPAR/carbohydrate.param
REMARK 3 PARAMETER FILE 3 : CNS_TOPPAR/water.param
REMARK 3 TOPOLOGY FILE 1 : CNS_TOPPAR/protein.top
30 REMARK 3 TOPOLOGY FILE 2 : NULL
REMARK 3
REMARK 3 OTHER REFINEMENT REMARKS: NULL
SEQRES 1 A 1440 VAL THR GLN ASP CYS LEU GLN LEU ILE ALA ASP SER GLU
SEQRES 2 A 1440 THR PRO THR ILE GLN LYS GLY SER TYR THR PHE VAL PRO
35 SEQRES 3 A 1440 TRP LEU LEU SER PHE LYS ARG GLY SER ALA LEU GLU GLU
SEQRES 4 A 1440 LYS GLU ASN LYS ILE LEU VAL LYS GLU THR GLY TYR PHE
SEQRES 5 A 1440 PHE ILE TYR GLY GLN VAL LEU TYR THR ASP LYS THR TYR
SEQRES 6 A 1440 ALA MET GLY HIS LEU ILE GLN ARG LYS LYS VAL HIS VAL
SEQRES 7 A 1440 PHE GLY ASP GLU LEU SER LEU VAL THR LEU PHE ARG CYS
40 SEQRES 8 A 1440 ILE GLN ASN MET PRO GLU THR LEU PRO ASN ASN SER CYS
SEQRES 9 A 1440 TYR SER ALA GLY ILE ALA LYS LEU GLU GLU GLY ASP GLU
SEQRES 10 A 1440 LEU GLN LEU ALA ILE PRO ARG GLU ASN ALA GLN ILE SER
SEQRES 11 A 1440 LEU ASP GLY ASP VAL THR PHE PHE GLY ALA LEU LYS LEU
SEQRES 12 A 1440 LEU VAL THR GLN ASP CYS LEU GLN LEU ILE ALA ASP SER
45 SEQRES 13 A 1440 GLU THR PRO THR ILE GLN LYS GLY SER TYR THR PHE VAL
SEQRES 14 A 1440 PRO TRP LEU LEU SER PHE LYS ARG GLY SER ALA LEU GLU
SEQRES 15 A 1440 GLU LYS GLU ASN LYS ILE LEU VAL LYS GLU THR GLY TYR
SEQRES 16 A 1440 PHE PHE ILE TYR GLY GLN VAL LEU TYR THR ASP LYS THR
SEQRES 17 A 1440 TYR ALA MET GLY HIS LEU ILE GLN ARG LYS LYS VAL HIS
50 SEQRES 18 A 1440 VAL PHE GLY ASP GLU LEU SER LEU VAL THR LEU PHE ARG
SEQRES 19 A 1440 CYS ILE GLN ASN MET PRO GLU THR LEU PRO ASN ASN SER
SEQRES 20 A 1440 CYS TYR SER ALA GLY ILE ALA LYS LEU GLU GLU GLY ASP
SEQRES 21 A 1440 GLU LEU GLN LEU ALA ILE PRO ARG GLU ASN ALA GLN ILE
SEQRES 22 A 1440 SER LEU ASP GLY ASP VAL THR PHE PHE GLY ALA LEU LYS
55 SEQRES 23 A 1440 LEU LEU VAL THR GLN ASP CYS LEU GLN LEU ILE ALA ASP
SEQRES 24 A 1440 SER GLU THR PRO THR ILE GLN LYS GLY SER TYR THR PHE
SEQRES 25 A 1440 VAL PRO TRP LEU LEU SER PHE LYS ARG GLY SER ALA LEU
SEQRES 26 A 1440 GLU GLU LYS GLU ASN LYS ILE LEU VAL LYS GLU THR GLY
SEQRES 27 A 1440 TYR PHE PHE ILE TYR GLY GLN VAL LEU TYR THR ASP LYS
60 SEQRES 28 A 1440 THR TYR ALA MET GLY HIS LEU ILE GLN ARG LYS LYS VAL
SEQRES 29 A 1440 HIS VAL PHE GLY ASP GLU LEU SER LEU VAL THR LEU PHE
SEQRES 30 A 1440 ARG CYS ILE GLN ASN MET PRO GLU THR LEU PRO ASN ASN
SEQRES 31 A 1440 SER CYS TYR SER ALA GLY ILE ALA LYS LEU GLU GLU GLY
SEQRES 32 A 1440 ASP GLU LEU GLN LEU ALA ILE PRO ARG GLU ASN ALA GLN
65 SEQRES 33 A 1440 ILE SER LEU ASP GLY ASP VAL THR PHE PHE GLY ALA LEU
SEQRES 34 A 1440 LYS LEU LEU VAL THR GLN ASP CYS LEU GLN LEU ILE ALA

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	SEQRES	35	A	1440	ASP	SER	GLU	THR	PRO	THR	ILE	GLN	LYS	GLY	SER	TYR	THR
	SEQRES	36	A	1440	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE	LYS	ARG	GLY	SER	ALA
	SEQRES	37	A	1440	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE	LEU	VAL	LYS	GLU	THR
	SEQRES	38	A	1440	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN	VAL	LEU	TYR	THR	ASP
5	SEQRES	39	A	1440	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU	ILE	GLN	ARG	LYS	LYS
	SEQRES	40	A	1440	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU	SER	LEU	VAL	THR	LEU
	SEQRES	41	A	1440	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO	GLU	THR	LEU	PRO	ASN
	SEQRES	42	A	1440	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE	ALA	LYS	LEU	GLU	GLU
	SEQRES	43	A	1440	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE	PRO	ARG	GLU	ASN	ALA
10	SEQRES	44	A	1440	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL	THR	PHE	PHE	GLY	ALA
	SEQRES	45	A	1440	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP	CYS	LEU	GLN	LEU	ILE
	SEQRES	46	A	1440	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	GLN	LYS	GLY	SER	TYR
	SEQRES	47	A	1440	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE	LYS	ARG	GLY	SER
	SEQRES	48	A	1440	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE	LEU	VAL	LYS	GLU
15	SEQRES	49	A	1440	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN	VAL	LEU	TYR	THR
	SEQRES	50	A	1440	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU	ILE	GLN	ARG	LYS
	SEQRES	51	A	1440	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU	SER	LEU	VAL	THR
	SEQRES	52	A	1440	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO	GLU	THR	LEU	PRO
	SEQRES	53	A	1440	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE	ALA	LYS	LEU	GLU
20	SEQRES	54	A	1440	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE	PRO	ARG	GLU	ASN
	SEQRES	55	A	1440	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL	THR	PHE	PHE	GLY
	SEQRES	56	A	1440	ALA	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP	CYS	LEU	GLN	LEU
	SEQRES	57	A	1440	ILE	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	GLN	LYS	GLY	SER
	SEQRES	58	A	1440	TYR	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE	LYS	ARG	GLY
25	SEQRES	59	A	1440	SER	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE	LEU	VAL	LYS
	SEQRES	60	A	1440	GLU	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN	VAL	LEU	TYR
	SEQRES	61	A	1440	THR	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU	ILE	GLN	ARG
	SEQRES	62	A	1440	LYS	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU	SER	LEU	VAL
	SEQRES	63	A	1440	THR	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO	GLU	THR	LEU
30	SEQRES	64	A	1440	PRO	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE	ALA	LYS	LEU
	SEQRES	65	A	1440	GLU	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE	PRO	ARG	GLU
	SEQRES	66	A	1440	ASN	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL	THR	PHE	PHE
	SEQRES	67	A	1440	GLY	ALA	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP	CYS	LEU	GLN
	SEQRES	68	A	1440	LEU	ILE	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	GLN	LYS	GLY
35	SEQRES	69	A	1440	SER	TYR	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE	LYS	ARG
	SEQRES	70	A	1440	GLY	SER	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE	LEU	VAL
	SEQRES	71	A	1440	LYS	GLU	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN	VAL	LEU
	SEQRES	72	A	1440	TYR	THR	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU	ILE	GLN
	SEQRES	73	A	1440	ARG	LYS	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU	SER	LEU
40	SEQRES	74	A	1440	VAL	THR	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO	GLU	THR
	SEQRES	75	A	1440	LEU	PRO	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE	ALA	LYS
	SEQRES	76	A	1440	LEU	GLU	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE	PRO	ARG
	SEQRES	77	A	1440	GLU	ASN	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL	THR	PHE
	SEQRES	78	A	1440	PHE	GLY	ALA	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP	CYS	LEU
45	SEQRES	79	A	1440	GLN	LEU	ILE	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	GLN	LYS
	SEQRES	80	A	1440	GLY	SER	TYR	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE	LYS
	SEQRES	81	A	1440	ARG	GLY	SER	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE	LEU
	SEQRES	82	A	1440	VAL	LYS	GLU	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN	VAL
	SEQRES	83	A	1440	LEU	TYR	THR	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU	ILE
50	SEQRES	84	A	1440	GLN	ARG	LYS	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU	SER
	SEQRES	85	A	1440	LEU	VAL	THR	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO	GLU
	SEQRES	86	A	1440	THR	LEU	PRO	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE	ALA
	SEQRES	87	A	1440	LYS	LEU	GLU	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE	PRO
	SEQRES	88	A	1440	ARG	GLU	ASN	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL	THR
55	SEQRES	89	A	1440	PHE	PHE	GLY	ALA	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP	CYS
	SEQRES	90	A	1440	LEU	GLN	LEU	ILE	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	GLN
	SEQRES	91	A	1440	LYS	GLY	SER	TYR	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	PHE
	SEQRES	92	A	1440	LYS	ARG	GLY	SER	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	ILE
	SEQRES	93	A	1440	LEU	VAL	LYS	GLU	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	GLN
60	SEQRES	94	A	1440	VAL	LEU	TYR	THR	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	LEU
	SEQRES	95	A	1440	ILE	GLN	ARG	LYS	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	LEU
	SEQRES	96	A	1440	SER	LEU	VAL	THR	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	PRO
	SEQRES	97	A	1440	GLU	THR	LEU	PRO	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	ILE
	SEQRES	98	A	1440	ALA	LYS	LEU	GLU	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	ILE
65	SEQRES	99	A	1440	PRO	ARG	GLU	ASN	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	VAL
	SEQRES	100	A	1440	THR	PHE	PHE	GLY	ALA	LEU	LYS	LEU	LEU	VAL	THR	GLN	ASP

	SEQRES	101	A	1440	CYS	LEU	GLN	LEU	ILE	ALA	ASP	SER	GLU	THR	PRO	THR	ILE	
	SEQRES	102	A	1440	GLN	LYS	GLY	SER	TYR	THR	PHE	VAL	PRO	TRP	LEU	LEU	SER	
	SEQRES	103	A	1440	PHE	LYS	ARG	GLY	SER	ALA	LEU	GLU	GLU	LYS	GLU	ASN	LYS	
	SEQRES	104	A	1440	ILE	LEU	VAL	LYS	GLU	THR	GLY	TYR	PHE	PHE	ILE	TYR	GLY	
5	SEQRES	105	A	1440	GLN	VAL	LEU	TYR	THR	ASP	LYS	THR	TYR	ALA	MET	GLY	HIS	
	SEQRES	106	A	1440	LEU	ILE	GLN	ARG	LYS	LYS	VAL	HIS	VAL	PHE	GLY	ASP	GLU	
	SEQRES	107	A	1440	LEU	SER	LEU	VAL	THR	LEU	PHE	ARG	CYS	ILE	GLN	ASN	MET	
	SEQRES	108	A	1440	PRO	GLU	THR	LEU	PRO	ASN	ASN	SER	CYS	TYR	SER	ALA	GLY	
	SEQRES	109	A	1440	ILE	ALA	LYS	LEU	GLU	GLU	GLY	ASP	GLU	LEU	GLN	LEU	ALA	
10	SEQRES	110	A	1440	ILE	PRO	ARG	GLU	ASN	ALA	GLN	ILE	SER	LEU	ASP	GLY	ASP	
	SEQRES	111	A	1440	VAL	THR	PHE	PHE	GLY	ALA	LEU	LYS	LEU	LEU				
	CRYST1	234.	242	234.	242	212.	550	90.00	90.00	120.00	P	63	2	2			12	
	ORIGX1			1.000000	0.000000	0.000000						0.000000						
	ORIGX2			0.000000	1.000000	0.000000						0.000000						
15	ORIGX3			0.000000	0.000000	1.000000						0.000000						
	SCALE1			0.004269	0.002465	0.000000						0.000000						
	SCALE2			0.000000	0.004930	0.000000						0.000000						
	SCALE3			0.000000	0.000000	0.004705						0.000000						
	ATOM	1	CB	VAL	A	1	-10.978	93.788	-41.518	1.00	25.10							T1
20	ATOM	2	CG1	VAL	A	1	-11.675	94.504	-40.346	1.00	31.97							T1
	ATOM	3	CG2	VAL	A	1	-10.397	94.817	-42.486	1.00	33.37							T1
	ATOM	4	C	VAL	A	1	-10.478	91.744	-40.116	1.00	32.82							T1
	ATOM	5	O	VAL	A	1	-10.182	91.658	-38.920	1.00	31.95							T1
	ATOM	6	N	VAL	A	1	-9.048	92.268	-42.107	1.00	31.73							T1
25	ATOM	7	CA	VAL	A	1	-9.842	92.845	-40.988	1.00	26.42							T1
	ATOM	8	N	THR	A	2	-11.344	90.908	-40.695	1.00	31.16							T1
	ATOM	9	CA	THR	A	2	-11.992	89.850	-39.916	1.00	36.43							T1
	ATOM	10	CB	THR	A	2	-13.529	89.918	-40.022	1.00	16.96							T1
	ATOM	11	OG1	THR	A	2	-13.937	89.499	-41.330	1.00	26.43							T1
30	ATOM	12	CG2	THR	A	2	-14.025	91.335	-39.751	1.00	31.50							T1
	ATOM	13	C	THR	A	2	-11.567	88.450	-40.338	1.00	26.63							T1
	ATOM	14	O	THR	A	2	-10.892	88.279	-41.351	1.00	30.53							T1
	ATOM	15	N	GLN	A	3	-11.981	87.452	-39.558	1.00	32.43							T1
	ATOM	16	CA	GLN	A	3	-11.649	86.053	-39.839	1.00	28.32							T1
35	ATOM	17	CB	GLN	A	3	-11.027	85.401	-38.605	1.00	28.94							T1
	ATOM	18	CG	GLN	A	3	-9.960	86.233	-37.940	1.00	34.72							T1
	ATOM	19	CD	GLN	A	3	-9.268	85.475	-36.834	1.00	31.04							T1
	ATOM	20	OE1	GLN	A	3	-8.658	84.435	-37.079	1.00	22.78							T1
	ATOM	21	NE2	GLN	A	3	-9.357	85.984	-35.608	1.00	33.37							T1
40	ATOM	22	C	GLN	A	3	-12.877	85.239	-40.248	1.00	25.83							T1
	ATOM	23	O	GLN	A	3	-13.689	84.863	-39.395	1.00	28.27							T1
	ATOM	24	N	ASP	A	4	-13.017	84.956	-41.538	1.00	32.26							T1
	ATOM	25	CA	ASP	A	4	-14.156	84.174	-41.982	1.00	29.72							T1
	ATOM	26	CB	ASP	A	4	-14.143	84.021	-43.496	1.00	32.34							T1
45	ATOM	27	CG	ASP	A	4	-14.289	85.344	-44.210	1.00	23.55							T1
	ATOM	28	OD1	ASP	A	4	-14.721	86.317	-43.550	1.00	26.88							T1
	ATOM	29	OD2	ASP	A	4	-13.987	85.414	-45.426	1.00	33.89							T1
	ATOM	30	C	ASP	A	4	-14.131	82.798	-41.344	1.00	24.29							T1
	ATOM	31	O	ASP	A	4	-13.063	82.246	-41.092	1.00	31.95							T1
50	ATOM	32	N	CYS	A	5	-15.312	82.252	-41.076	1.00	34.21							T1
	ATOM	33	CA	CYS	A	5	-15.446	80.930	-40.475	1.00	32.35							T1
	ATOM	34	CB	CYS	A	5	-15.273	80.996	-38.939	1.00	21.67							T1
	ATOM	35	SG	CYS	A	5	-15.771	82.553	-38.115	1.00	30.78							T1
	ATOM	36	C	CYS	A	5	-16.800	80.338	-40.822	1.00	28.48							T1
55	ATOM	37	O	CYS	A	5	-17.771	81.056	-40.985	1.00	31.54							T1
	ATOM	38	N	LEU	A	6	-16.852	79.023	-40.967	1.00	29.50							T1
	ATOM	39	CA	LEU	A	6	-18.103	78.335	-41.273	1.00	34.06							T1
	ATOM	40	CB	LEU	A	6	-18.178	77.982	-42.758	1.00	30.50							T1
	ATOM	41	CG	LEU	A	6	-19.415	77.203	-43.224	1.00	26.08							T1
60	ATOM	42	CD1	LEU	A	6	-19.719	77.543	-44.654	1.00	24.45							T1
	ATOM	43	CD2	LEU	A	6	-19.183	75.718	-43.077	1.00	33.62							T1
	ATOM	44	C	LEU	A	6	-18.143	77.072	-40.426	1.00	33.39							T1
	ATOM	45	O	LEU	A	6	-17.157	76.348	-40.349	1.00	33.50							T1
	ATOM	46	N	GLN	A	7	-19.276	76.812	-39.784	1.00	34.98							T1
65	ATOM	47	CA	GLN	A	7	-19.395	75.642	-38.931	1.00	26.98							T1
	ATOM	48	CB	GLN	A	7	-19.436	76.071	-37.474	1.00	31.79							T1

	ATOM	49	CG	GLN	A	7	-19.283	74.939	-36.491	1.00	33.29	T1
	ATOM	50	CD	GLN	A	7	-19.084	75.454	-35.092	1.00	29.25	T1
	ATOM	51	OE1	GLN	A	7	-19.991	76.022	-34.499	1.00	30.48	T1
	ATOM	52	NE2	GLN	A	7	-17.888	75.283	-34.562	1.00	29.89	T1
5	ATOM	53	C	GLN	A	7	-20.624	74.823	-39.246	1.00	26.97	T1
	ATOM	54	O	GLN	A	7	-21.687	75.369	-39.522	1.00	33.60	T1
	ATOM	55	N	LEU	A	8	-20.467	73.506	-39.198	1.00	31.83	T1
	ATOM	56	CA	LEU	A	8	-21.557	72.586	-39.484	1.00	24.41	T1
	ATOM	57	CB	LEU	A	8	-21.175	71.657	-40.641	1.00	38.54	T1
10	ATOM	58	CG	LEU	A	8	-21.404	72.087	-42.095	1.00	36.96	T1
	ATOM	59	CD1	LEU	A	8	-21.669	73.566	-42.196	1.00	32.34	T1
	ATOM	60	CD2	LEU	A	8	-20.199	71.696	-42.912	1.00	29.78	T1
	ATOM	61	C	LEU	A	8	-21.928	71.758	-38.263	1.00	31.29	T1
	ATOM	62	O	LEU	A	8	-21.098	71.509	-37.394	1.00	28.94	T1
15	ATOM	63	N	ILE	A	9	-23.185	71.328	-38.222	1.00	31.60	T1
	ATOM	64	CA	ILE	A	9	-23.730	70.530	-37.130	1.00	31.98	T1
	ATOM	65	CB	ILE	A	9	-24.781	71.330	-36.362	1.00	39.47	T1
	ATOM	66	CG2	ILE	A	9	-25.652	70.413	-35.558	1.00	28.69	T1
	ATOM	67	CG1	ILE	A	9	-24.123	72.327	-35.433	1.00	36.23	T1
20	ATOM	68	CD1	ILE	A	9	-25.154	73.150	-34.702	1.00	28.20	T1
	ATOM	69	C	ILE	A	9	-24.424	69.298	-37.698	1.00	26.36	T1
	ATOM	70	O	ILE	A	9	-25.053	69.365	-38.751	1.00	35.46	T1
	ATOM	71	N	ALA	A	10	-24.335	68.178	-36.992	1.00	29.29	T1
	ATOM	72	CA	ALA	A	10	-24.996	66.960	-37.459	1.00	26.96	T1
25	ATOM	73	CB	ALA	A	10	-24.701	65.799	-36.527	1.00	25.44	T1
	ATOM	74	C	ALA	A	10	-26.504	67.163	-37.552	1.00	22.64	T1
	ATOM	75	O	ALA	A	10	-27.141	67.649	-36.608	1.00	25.37	T1
	ATOM	76	N	ASP	A	11	-27.073	66.789	-38.695	1.00	28.47	T1
	ATOM	77	CA	ASP	A	11	-28.513	66.910	-38.903	1.00	24.77	T1
30	ATOM	78	CB	ASP	A	11	-28.828	67.197	-40.369	1.00	37.13	T1
	ATOM	79	CG	ASP	A	11	-30.307	67.199	-40.641	1.00	31.60	T1
	ATOM	80	OD1	ASP	A	11	-31.054	67.693	-39.766	1.00	30.74	T1
	ATOM	81	OD2	ASP	A	11	-30.719	66.719	-41.720	1.00	32.60	T1
	ATOM	82	C	ASP	A	11	-29.200	65.625	-38.471	1.00	35.17	T1
35	ATOM	83	O	ASP	A	11	-29.370	64.699	-39.266	1.00	23.07	T1
	ATOM	84	N	SER	A	12	-29.581	65.595	-37.196	1.00	32.64	T1
	ATOM	85	CA	SER	A	12	-30.237	64.450	-36.578	1.00	35.92	T1
	ATOM	86	CB	SER	A	12	-30.407	64.697	-35.070	1.00	29.42	T1
	ATOM	87	OG	SER	A	12	-31.174	65.871	-34.810	1.00	27.48	T1
40	ATOM	88	C	SER	A	12	-31.599	64.117	-37.184	1.00	29.05	T1
	ATOM	89	O	SER	A	12	-32.431	63.485	-36.538	1.00	38.27	T1
	ATOM	90	N	GLU	A	13	-31.835	64.541	-38.418	1.00	27.32	T1
	ATOM	91	CA	GLU	A	13	-33.114	64.251	-39.042	1.00	22.33	T1
	ATOM	92	CB	GLU	A	13	-34.037	65.446	-38.921	1.00	31.69	T1
45	ATOM	93	CG	GLU	A	13	-34.844	65.384	-37.656	1.00	35.73	T1
	ATOM	94	CD	GLU	A	13	-35.842	66.513	-37.574	1.00	29.70	T1
	ATOM	95	OE1	GLU	A	13	-36.475	66.821	-38.617	1.00	28.48	T1
	ATOM	96	OE2	GLU	A	13	-36.002	67.097	-36.471	1.00	31.34	T1
	ATOM	97	C	GLU	A	13	-33.070	63.770	-40.479	1.00	34.29	T1
50	ATOM	98	O	GLU	A	13	-34.034	63.937	-41.230	1.00	32.43	T1
	ATOM	99	N	THR	A	14	-31.941	63.184	-40.857	1.00	32.83	T1
	ATOM	100	CA	THR	A	14	-31.778	62.614	-42.180	1.00	26.15	T1
	ATOM	101	CB	THR	A	14	-31.200	63.608	-43.210	1.00	32.96	T1
	ATOM	102	OG1	THR	A	14	-29.945	64.092	-42.742	1.00	22.86	T1
55	ATOM	103	CG2	THR	A	14	-32.149	64.779	-43.432	1.00	26.33	T1
	ATOM	104	C	THR	A	14	-30.823	61.449	-41.987	1.00	33.69	T1
	ATOM	105	O	THR	A	14	-29.983	61.443	-41.077	1.00	33.30	T1
	ATOM	106	N	PRO	A	15	-30.959	60.425	-42.831	1.00	28.41	T1
	ATOM	107	CD	PRO	A	15	-31.910	60.331	-43.950	1.00	30.43	T1
60	ATOM	108	CA	PRO	A	15	-30.111	59.231	-42.758	1.00	35.28	T1
	ATOM	109	CB	PRO	A	15	-30.668	58.341	-43.875	1.00	26.85	T1
	ATOM	110	CG	PRO	A	15	-32.086	58.847	-44.064	1.00	35.62	T1
	ATOM	111	C	PRO	A	15	-28.649	59.573	-42.997	1.00	25.22	T1
	ATOM	112	O	PRO	A	15	-28.332	60.461	-43.788	1.00	23.17	T1
65	ATOM	113	N	THR	A	16	-27.761	58.866	-42.322	1.00	31.09	T1
	ATOM	114	CA	THR	A	16	-26.344	59.105	-42.515	1.00	34.31	T1

	ATOM	115	CB	THR	A	16	-25.540	58.367	-41.455	1.00	30.03	T1
	ATOM	116	OG1	THR	A	16	-25.607	56.954	-41.692	1.00	23.86	T1
	ATOM	117	CG2	THR	A	16	-26.120	58.655	-40.087	1.00	30.14	T1
	ATOM	118	C	THR	A	16	-25.954	58.584	-43.899	1.00	27.65	T1
5	ATOM	119	O	THR	A	16	-26.050	57.383	-44.169	1.00	30.95	T1
	ATOM	120	N	ILE	A	17	-25.516	59.484	-44.774	1.00	30.99	T1
	ATOM	121	CA	ILE	A	17	-25.128	59.107	-46.136	1.00	27.33	T1
	ATOM	122	CB	ILE	A	17	-24.536	60.309	-46.883	1.00	28.06	T1
	ATOM	123	CG2	ILE	A	17	-24.125	59.902	-48.278	1.00	31.47	T1
10	ATOM	124	CG1	ILE	A	17	-25.568	61.433	-46.939	1.00	27.05	T1
	ATOM	125	CD1	ILE	A	17	-25.061	62.694	-47.590	1.00	34.82	T1
	ATOM	126	C	ILE	A	17	-24.134	57.940	-46.238	1.00	29.81	T1
	ATOM	127	O	ILE	A	17	-23.110	57.913	-45.549	1.00	24.15	T1
	ATOM	128	N	GLN	A	18	-24.454	56.978	-47.105	1.00	28.69	T1
15	ATOM	129	CA	GLN	A	18	-23.603	55.813	-47.325	1.00	34.08	T1
	ATOM	130	CB	GLN	A	18	-24.384	54.524	-47.115	1.00	26.34	T1
	ATOM	131	CG	GLN	A	18	-23.744	53.624	-46.097	1.00	28.43	T1
	ATOM	132	CD	GLN	A	18	-23.927	54.165	-44.703	1.00	31.90	T1
	ATOM	133	OE1	GLN	A	18	-25.010	54.054	-44.119	1.00	24.64	T1
20	ATOM	134	NE2	GLN	A	18	-22.877	54.777	-44.162	1.00	29.89	T1
	ATOM	135	C	GLN	A	18	-23.043	55.818	-48.740	1.00	25.18	T1
	ATOM	136	O	GLN	A	18	-23.774	56.057	-49.704	1.00	29.43	T1
	ATOM	137	N	LYS	A	19	-21.750	55.536	-48.869	1.00	35.16	T1
	ATOM	138	CA	LYS	A	19	-21.118	55.533	-50.181	1.00	33.75	T1
25	ATOM	139	CB	LYS	A	19	-21.141	56.947	-50.766	1.00	33.23	T1
	ATOM	140	CG	LYS	A	19	-20.293	57.119	-52.013	1.00	29.08	T1
	ATOM	141	CD	LYS	A	19	-20.521	58.486	-52.643	1.00	30.83	T1
	ATOM	142	CE	LYS	A	19	-19.648	58.681	-53.888	1.00	37.17	T1
	ATOM	143	NZ	LYS	A	19	-19.899	59.989	-54.580	1.00	29.78	T1
30	ATOM	144	C	LYS	A	19	-19.687	55.016	-50.142	1.00	29.76	T1
	ATOM	145	O	LYS	A	19	-18.875	55.456	-49.317	1.00	31.02	T1
	ATOM	146	N	GLY	A	20	-19.382	54.091	-51.052	1.00	29.61	T1
	ATOM	147	CA	GLY	A	20	-18.048	53.514	-51.115	1.00	31.01	T1
	ATOM	148	C	GLY	A	20	-17.658	52.950	-49.763	1.00	26.54	T1
35	ATOM	149	O	GLY	A	20	-16.516	53.112	-49.326	1.00	33.31	T1
	ATOM	150	N	SER	A	21	-18.617	52.297	-49.103	1.00	22.69	T1
	ATOM	151	CA	SER	A	21	-18.408	51.700	-47.780	1.00	35.68	T1
	ATOM	152	CB	SER	A	21	-17.529	50.440	-47.894	1.00	29.04	T1
	ATOM	153	OG	SER	A	21	-16.246	50.729	-48.429	1.00	28.47	T1
40	ATOM	154	C	SER	A	21	-17.804	52.695	-46.773	1.00	30.82	T1
	ATOM	155	O	SER	A	21	-16.940	52.353	-45.952	1.00	38.73	T1
	ATOM	156	N	TYR	A	22	-18.284	53.931	-46.863	1.00	34.91	T1
	ATOM	157	CA	TYR	A	22	-17.872	55.025	-45.989	1.00	32.43	T1
	ATOM	158	CB	TYR	A	22	-17.104	56.080	-46.787	1.00	27.02	T1
45	ATOM	159	CG	TYR	A	22	-15.608	55.937	-46.738	1.00	35.57	T1
	ATOM	160	CD1	TYR	A	22	-15.014	54.707	-46.479	1.00	28.37	T1
	ATOM	161	CE1	TYR	A	22	-13.623	54.567	-46.465	1.00	24.81	T1
	ATOM	162	CD2	TYR	A	22	-14.778	57.032	-46.984	1.00	34.83	T1
	ATOM	163	CE2	TYR	A	22	-13.387	56.904	-46.976	1.00	25.19	T1
50	ATOM	164	CZ	TYR	A	22	-12.818	55.668	-46.715	1.00	27.09	T1
	ATOM	165	OH	TYR	A	22	-11.450	55.525	-46.702	1.00	33.19	T1
	ATOM	166	C	TYR	A	22	-19.153	55.650	-45.451	1.00	28.19	T1
	ATOM	167	O	TYR	A	22	-20.172	55.672	-46.146	1.00	30.21	T1
	ATOM	168	N	THR	A	23	-19.118	56.147	-44.222	1.00	31.15	T1
55	ATOM	169	CA	THR	A	23	-20.301	56.785	-43.680	1.00	30.35	T1
	ATOM	170	CB	THR	A	23	-20.620	56.283	-42.273	1.00	31.36	T1
	ATOM	171	OG1	THR	A	23	-20.408	54.867	-42.206	1.00	31.37	T1
	ATOM	172	CG2	THR	A	23	-22.072	56.573	-41.943	1.00	29.36	T1
	ATOM	173	C	THR	A	23	-20.051	58.285	-43.631	1.00	28.17	T1
60	ATOM	174	O	THR	A	23	-19.000	58.734	-43.164	1.00	24.88	T1
	ATOM	175	N	PHE	A	24	-21.006	59.056	-44.136	1.00	31.90	T1
	ATOM	176	CA	PHE	A	24	-20.883	60.504	-44.134	1.00	31.66	T1
	ATOM	177	CB	PHE	A	24	-20.916	61.038	-45.557	1.00	35.94	T1
	ATOM	178	CG	PHE	A	24	-19.737	60.622	-46.380	1.00	25.35	T1
65	ATOM	179	CD1	PHE	A	24	-19.683	59.356	-46.957	1.00	33.16	T1
	ATOM	180	CD2	PHE	A	24	-18.661	61.492	-46.566	1.00	35.82	T1

	ATOM	181	CE1	PHE	A	24	-18.571	58.966	-47.710	1.00	23.65	T1
	ATOM	182	CE2	PHE	A	24	-17.550	61.111	-47.312	1.00	35.78	T1
	ATOM	183	CZ	PHE	A	24	-17.505	59.848	-47.884	1.00	40.18	T1
	ATOM	184	C	PHE	A	24	-21.980	61.169	-43.317	1.00	35.66	T1
5	ATOM	185	O	PHE	A	24	-23.167	60.958	-43.560	1.00	25.47	T1
	ATOM	186	N	VAL	A	25	-21.571	61.974	-42.343	1.00	32.22	T1
	ATOM	187	CA	VAL	A	25	-22.507	62.680	-41.477	1.00	27.17	T1
	ATOM	188	CB	VAL	A	25	-21.765	63.437	-40.360	1.00	27.67	T1
	ATOM	189	CG1	VAL	A	25	-22.744	64.232	-39.527	1.00	35.16	T1
10	ATOM	190	CG2	VAL	A	25	-20.992	62.463	-39.503	1.00	35.87	T1
	ATOM	191	C	VAL	A	25	-23.325	63.697	-42.266	1.00	31.85	T1
	ATOM	192	O	VAL	A	25	-22.781	64.457	-43.070	1.00	25.37	T1
	ATOM	193	N	PRO	A	26	-24.651	63.710	-42.056	1.00	39.81	T1
	ATOM	194	CD	PRO	A	26	-25.421	62.723	-41.279	1.00	25.78	T1
15	ATOM	195	CA	PRO	A	26	-25.549	64.648	-42.746	1.00	27.02	T1
	ATOM	196	CB	PRO	A	26	-26.936	64.071	-42.472	1.00	31.01	T1
	ATOM	197	CG	PRO	A	26	-26.670	62.630	-42.081	1.00	27.89	T1
	ATOM	198	C	PRO	A	26	-25.376	66.007	-42.066	1.00	30.66	T1
	ATOM	199	O	PRO	A	26	-25.767	66.169	-40.910	1.00	29.07	T1
20	ATOM	200	N	TRP	A	27	-24.800	66.980	-42.762	1.00	39.09	T1
	ATOM	201	CA	TRP	A	27	-24.591	68.280	-42.146	1.00	30.23	T1
	ATOM	202	CB	TRP	A	27	-23.343	68.944	-42.724	1.00	38.71	T1
	ATOM	203	CG	TRP	A	27	-22.097	68.187	-42.432	1.00	28.02	T1
	ATOM	204	CD2	TRP	A	27	-21.626	67.764	-41.145	1.00	29.26	T1
25	ATOM	205	CE2	TRP	A	27	-20.442	67.030	-41.355	1.00	26.91	T1
	ATOM	206	CE3	TRP	A	27	-22.094	67.931	-39.836	1.00	28.91	T1
	ATOM	207	CD1	TRP	A	27	-21.203	67.714	-43.341	1.00	28.87	T1
	ATOM	208	NE1	TRP	A	27	-20.205	67.015	-42.704	1.00	31.75	T1
	ATOM	209	CZ2	TRP	A	27	-19.715	66.462	-40.307	1.00	28.21	T1
30	ATOM	210	CZ3	TRP	A	27	-21.370	67.365	-38.792	1.00	30.61	T1
	ATOM	211	CH2	TRP	A	27	-20.194	66.638	-39.037	1.00	33.04	T1
	ATOM	212	C	TRP	A	27	-25.751	69.253	-42.217	1.00	28.49	T1
	ATOM	213	O	TRP	A	27	-26.687	69.099	-42.998	1.00	34.59	T1
	ATOM	214	N	LEU	A	28	-25.658	70.269	-41.372	1.00	31.56	T1
35	ATOM	215	CA	LEU	A	28	-26.653	71.321	-41.283	1.00	28.42	T1
	ATOM	216	CB	LEU	A	28	-27.705	70.950	-40.243	1.00	34.42	T1
	ATOM	217	CG	LEU	A	28	-29.021	71.708	-40.363	1.00	28.77	T1
	ATOM	218	CD1	LEU	A	28	-29.666	71.369	-41.724	1.00	38.00	T1
	ATOM	219	CD2	LEU	A	28	-29.932	71.324	-39.197	1.00	39.34	T1
40	ATOM	220	C	LEU	A	28	-25.896	72.567	-40.838	1.00	22.38	T1
	ATOM	221	O	LEU	A	28	-25.158	72.533	-39.851	1.00	35.84	T1
	ATOM	222	N	LEU	A	29	-26.072	73.665	-41.559	1.00	26.93	T1
	ATOM	223	CA	LEU	A	29	-25.361	74.886	-41.212	1.00	28.00	T1
	ATOM	224	CB	LEU	A	29	-25.776	76.036	-42.126	1.00	30.08	T1
45	ATOM	225	CG	LEU	A	29	-25.067	77.343	-41.784	1.00	30.64	T1
	ATOM	226	CD1	LEU	A	29	-23.642	77.283	-42.289	1.00	31.54	T1
	ATOM	227	CD2	LEU	A	29	-25.790	78.502	-42.411	1.00	21.43	T1
	ATOM	228	C	LEU	A	29	-25.572	75.324	-39.769	1.00	26.94	T1
	ATOM	229	O	LEU	A	29	-26.706	75.475	-39.306	1.00	37.48	T1
50	ATOM	230	N	SER	A	30	-24.467	75.517	-39.060	1.00	31.38	T1
	ATOM	231	CA	SER	A	30	-24.526	75.987	-37.691	1.00	27.56	T1
	ATOM	232	CB	SER	A	30	-23.307	75.528	-36.902	1.00	32.55	T1
	ATOM	233	OG	SER	A	30	-23.328	76.095	-35.605	1.00	31.47	T1
	ATOM	234	C	SER	A	30	-24.503	77.497	-37.826	1.00	33.12	T1
55	ATOM	235	O	SER	A	30	-25.378	78.188	-37.328	1.00	23.94	T1
	ATOM	236	N	PHE	A	31	-23.495	78.003	-38.521	1.00	26.59	T1
	ATOM	237	CA	PHE	A	31	-23.371	79.435	-38.751	1.00	17.90	T1
	ATOM	238	CB	PHE	A	31	-23.003	80.160	-37.451	1.00	27.14	T1
	ATOM	239	CG	PHE	A	31	-21.533	80.217	-37.180	1.00	27.03	T1
60	ATOM	240	CD1	PHE	A	31	-20.745	81.204	-37.759	1.00	33.93	T1
	ATOM	241	CD2	PHE	A	31	-20.931	79.274	-36.356	1.00	24.60	T1
	ATOM	242	CE1	PHE	A	31	-19.379	81.252	-37.524	1.00	27.40	T1
	ATOM	243	CE2	PHE	A	31	-19.564	79.313	-36.113	1.00	31.50	T1
	ATOM	244	CZ	PHE	A	31	-18.785	80.306	-36.700	1.00	31.45	T1
65	ATOM	245	C	PHE	A	31	-22.302	79.664	-39.813	1.00	25.17	T1
	ATOM	246	O	PHE	A	31	-21.401	78.848	-39.986	1.00	31.10	T1

	ATOM	247	N	LYS	A	32	-22.419	80.767	-40.535	1.00	29.35	T1
	ATOM	248	CA	LYS	A	32	-21.465	81.100	-41.577	1.00	34.81	T1
	ATOM	249	CB	LYS	A	32	-22.064	80.833	-42.952	1.00	28.69	T1
	ATOM	250	CG	LYS	A	32	-21.249	81.427	-44.056	1.00	28.01	T1
5	ATOM	251	CD	LYS	A	32	-21.949	81.367	-45.382	1.00	35.83	T1
	ATOM	252	CE	LYS	A	32	-21.195	82.225	-46.377	1.00	37.12	T1
	ATOM	253	NZ	LYS	A	32	-21.734	82.070	-47.742	1.00	22.67	T1
	ATOM	254	C	LYS	A	32	-21.102	82.569	-41.446	1.00	28.42	T1
	ATOM	255	O	LYS	A	32	-21.970	83.431	-41.417	1.00	33.58	T1
10	ATOM	256	N	ARG	A	33	-19.812	82.853	-41.368	1.00	35.63	T1
	ATOM	257	CA	ARG	A	33	-19.343	84.221	-41.201	1.00	37.28	T1
	ATOM	258	CB	ARG	A	33	-18.761	84.382	-39.789	1.00	32.78	T1
	ATOM	259	CG	ARG	A	33	-18.084	85.699	-39.480	1.00	26.57	T1
	ATOM	260	CD	ARG	A	33	-18.040	85.886	-37.967	1.00	34.16	T1
15	ATOM	261	NE	ARG	A	33	-17.315	87.087	-37.549	1.00	25.05	T1
	ATOM	262	CZ	ARG	A	33	-15.993	87.153	-37.440	1.00	20.10	T1
	ATOM	263	NH1	ARG	A	33	-15.255	86.080	-37.715	1.00	29.86	T1
	ATOM	264	NH2	ARG	A	33	-15.413	88.286	-37.069	1.00	38.85	T1
	ATOM	265	C	ARG	A	33	-18.302	84.549	-42.248	1.00	27.32	T1
20	ATOM	266	O	ARG	A	33	-17.291	83.863	-42.372	1.00	31.88	T1
	ATOM	267	N	GLY	A	34	-18.558	85.598	-43.015	1.00	32.07	T1
	ATOM	268	CA	GLY	A	34	-17.608	85.982	-44.040	1.00	27.12	T1
	ATOM	269	C	GLY	A	34	-17.903	85.375	-45.399	1.00	28.02	T1
	ATOM	270	O	GLY	A	34	-18.997	84.837	-45.641	1.00	30.64	T1
25	ATOM	271	N	SER	A	35	-16.915	85.437	-46.285	1.00	27.59	T1
	ATOM	272	CA	SER	A	35	-17.080	84.932	-47.638	1.00	31.43	T1
	ATOM	273	CB	SER	A	35	-16.904	86.082	-48.613	1.00	30.24	T1
	ATOM	274	OG	SER	A	35	-15.652	86.720	-48.386	1.00	27.09	T1
	ATOM	275	C	SER	A	35	-16.143	83.810	-48.055	1.00	31.30	T1
30	ATOM	276	O	SER	A	35	-16.434	83.093	-49.011	1.00	35.13	T1
	ATOM	277	N	ALA	A	36	-15.024	83.656	-47.355	1.00	31.03	T1
	ATOM	278	CA	ALA	A	36	-14.040	82.635	-47.710	1.00	32.64	T1
	ATOM	279	CB	ALA	A	36	-12.796	82.801	-46.854	1.00	33.38	T1
	ATOM	280	C	ALA	A	36	-14.503	81.178	-47.668	1.00	32.11	T1
35	ATOM	281	O	ALA	A	36	-13.844	80.310	-48.249	1.00	37.23	T1
	ATOM	282	N	LEU	A	37	-15.620	80.897	-46.998	1.00	32.12	T1
	ATOM	283	CA	LEU	A	37	-16.113	79.520	-46.909	1.00	28.93	T1
	ATOM	284	CB	LEU	A	37	-15.761	78.943	-45.537	1.00	26.42	T1
	ATOM	285	CG	LEU	A	37	-14.265	78.899	-45.240	1.00	33.01	T1
40	ATOM	286	CD1	LEU	A	37	-14.033	78.761	-43.756	1.00	29.55	T1
	ATOM	287	CD2	LEU	A	37	-13.638	77.757	-46.013	1.00	28.97	T1
	ATOM	288	C	LEU	A	37	-17.616	79.401	-47.158	1.00	30.74	T1
	ATOM	289	O	LEU	A	37	-18.384	80.305	-46.822	1.00	30.51	T1
	ATOM	290	N	GLU	A	38	-18.028	78.275	-47.738	1.00	25.28	T1
45	ATOM	291	CA	GLU	A	38	-19.437	78.025	-48.057	1.00	30.11	T1
	ATOM	292	CB	GLU	A	38	-19.730	78.443	-49.496	1.00	23.84	T1
	ATOM	293	CG	GLU	A	38	-19.997	79.912	-49.715	1.00	37.28	T1
	ATOM	294	CD	GLU	A	38	-20.021	80.263	-51.188	1.00	27.08	T1
	ATOM	295	OE1	GLU	A	38	-20.484	79.420	-52.004	1.00	37.53	T1
50	ATOM	296	OE2	GLU	A	38	-19.581	81.389	-51.525	1.00	25.19	T1
	ATOM	297	C	GLU	A	38	-19.817	76.558	-47.925	1.00	30.68	T1
	ATOM	298	O	GLU	A	38	-18.942	75.695	-47.876	1.00	33.89	T1
	ATOM	299	N	GLU	A	39	-21.120	76.276	-47.870	1.00	29.60	T1
	ATOM	300	CA	GLU	A	39	-21.575	74.889	-47.798	1.00	30.02	T1
55	ATOM	301	CB	GLU	A	39	-22.915	74.748	-47.112	1.00	32.49	T1
	ATOM	302	CG	GLU	A	39	-23.106	75.557	-45.882	1.00	31.04	T1
	ATOM	303	CD	GLU	A	39	-24.572	75.910	-45.697	1.00	25.66	T1
	ATOM	304	OE1	GLU	A	39	-24.961	77.043	-46.084	1.00	29.40	T1
	ATOM	305	OE2	GLU	A	39	-25.337	75.043	-45.191	1.00	36.68	T1
60	ATOM	306	C	GLU	A	39	-21.809	74.484	-49.228	1.00	27.35	T1
	ATOM	307	O	GLU	A	39	-22.223	75.307	-50.044	1.00	31.54	T1
	ATOM	308	N	LYS	A	40	-21.567	73.222	-49.539	1.00	35.33	T1
	ATOM	309	CA	LYS	A	40	-21.806	72.755	-50.890	1.00	35.83	T1
	ATOM	310	CB	LYS	A	40	-20.687	73.192	-51.833	1.00	25.63	T1
65	ATOM	311	CG	LYS	A	40	-20.928	72.762	-53.272	1.00	27.77	T1
	ATOM	312	CD	LYS	A	40	-19.665	72.872	-54.106	1.00	28.81	T1

	ATOM	313	CE	LYS	A	40	-19.856	72.233	-55.477	1.00	29.48	T1
	ATOM	314	NZ	LYS	A	40	-18.567	72.194	-56.235	1.00	32.25	T1
	ATOM	315	C	LYS	A	40	-21.921	71.250	-50.903	1.00	29.49	T1
	ATOM	316	O	LYS	A	40	-20.923	70.535	-50.798	1.00	35.78	T1
5	ATOM	317	N	GLU	A	41	-23.153	70.776	-51.008	1.00	35.24	T1
	ATOM	318	CA	GLU	A	41	-23.408	69.351	-51.054	1.00	34.91	T1
	ATOM	319	CB	GLU	A	41	-22.974	68.822	-52.412	1.00	32.66	T1
	ATOM	320	CG	GLU	A	41	-23.555	69.656	-53.542	1.00	27.42	T1
	ATOM	321	CD	GLU	A	41	-22.875	69.407	-54.879	1.00	34.13	T1
10	ATOM	322	OE1	GLU	A	41	-21.614	69.507	-54.951	1.00	29.43	T1
	ATOM	323	OE2	GLU	A	41	-23.607	69.120	-55.864	1.00	31.13	T1
	ATOM	324	C	GLU	A	41	-22.697	68.611	-49.926	1.00	28.00	T1
	ATOM	325	O	GLU	A	41	-21.863	67.728	-50.153	1.00	31.54	T1
	ATOM	326	N	ASN	A	42	-23.028	69.000	-48.703	1.00	38.83	T1
15	ATOM	327	CA	ASN	A	42	-22.477	68.375	-47.513	1.00	36.40	T1
	ATOM	328	CB	ASN	A	42	-22.864	66.907	-47.475	1.00	29.03	T1
	ATOM	329	CG	ASN	A	42	-23.294	66.474	-46.103	1.00	26.93	T1
	ATOM	330	OD1	ASN	A	42	-22.776	65.502	-45.567	1.00	28.29	T1
	ATOM	331	ND2	ASN	A	42	-24.250	67.196	-45.519	1.00	26.56	T1
20	ATOM	332	C	ASN	A	42	-20.976	68.488	-47.345	1.00	33.46	T1
	ATOM	333	O	ASN	A	42	-20.374	67.706	-46.607	1.00	35.13	T1
	ATOM	334	N	LYS	A	43	-20.377	69.458	-48.027	1.00	24.57	T1
	ATOM	335	CA	LYS	A	43	-18.944	69.686	-47.934	1.00	33.62	T1
	ATOM	336	CB	LYS	A	43	-18.249	69.198	-49.199	1.00	26.05	T1
25	ATOM	337	CG	LYS	A	43	-18.253	67.697	-49.357	1.00	30.19	T1
	ATOM	338	CD	LYS	A	43	-17.598	67.282	-50.656	1.00	27.53	T1
	ATOM	339	CE	LYS	A	43	-18.480	67.638	-51.842	1.00	25.17	T1
	ATOM	340	NZ	LYS	A	43	-17.882	67.207	-53.149	1.00	30.10	T1
	ATOM	341	C	LYS	A	43	-18.686	71.169	-47.751	1.00	27.58	T1
30	ATOM	342	O	LYS	A	43	-19.548	71.993	-48.035	1.00	32.44	T1
	ATOM	343	N	ILE	A	44	-17.504	71.511	-47.262	1.00	29.27	T1
	ATOM	344	CA	ILE	A	44	-17.170	72.908	-47.082	1.00	28.78	T1
	ATOM	345	CB	ILE	A	44	-16.348	73.126	-45.815	1.00	27.76	T1
	ATOM	346	CG2	ILE	A	44	-16.013	74.597	-45.678	1.00	27.14	T1
35	ATOM	347	CG1	ILE	A	44	-17.135	72.646	-44.599	1.00	31.59	T1
	ATOM	348	CD1	ILE	A	44	-16.419	72.812	-43.304	1.00	29.67	T1
	ATOM	349	C	ILE	A	44	-16.354	73.347	-48.290	1.00	29.76	T1
	ATOM	350	O	ILE	A	44	-15.325	72.756	-48.603	1.00	24.71	T1
	ATOM	351	N	LEU	A	45	-16.822	74.382	-48.972	1.00	24.88	T1
40	ATOM	352	CA	LEU	A	45	-16.144	74.883	-50.155	1.00	31.33	T1
	ATOM	353	CB	LEU	A	45	-17.177	75.243	-51.216	1.00	37.65	T1
	ATOM	354	CG	LEU	A	45	-16.584	75.858	-52.480	1.00	25.53	T1
	ATOM	355	CD1	LEU	A	45	-15.820	74.794	-53.256	1.00	37.76	T1
	ATOM	356	CD2	LEU	A	45	-17.694	76.451	-53.311	1.00	33.03	T1
45	ATOM	357	C	LEU	A	45	-15.264	76.098	-49.879	1.00	31.30	T1
	ATOM	358	O	LEU	A	45	-15.707	77.081	-49.282	1.00	32.64	T1
	ATOM	359	N	VAL	A	46	-14.018	76.031	-50.332	1.00	33.56	T1
	ATOM	360	CA	VAL	A	46	-13.082	77.130	-50.135	1.00	29.47	T1
	ATOM	361	CB	VAL	A	46	-11.637	76.617	-50.129	1.00	23.29	T1
50	ATOM	362	CG1	VAL	A	46	-10.677	77.777	-49.929	1.00	36.02	T1
	ATOM	363	CG2	VAL	A	46	-11.463	75.601	-49.027	1.00	31.15	T1
	ATOM	364	C	VAL	A	46	-13.228	78.185	-51.230	1.00	22.73	T1
	ATOM	365	O	VAL	A	46	-13.043	77.893	-52.406	1.00	25.54	T1
	ATOM	366	N	LYS	A	47	-13.547	79.413	-50.845	1.00	30.60	T1
55	ATOM	367	CA	LYS	A	47	-13.727	80.478	-51.820	1.00	27.24	T1
	ATOM	368	CB	LYS	A	47	-15.001	81.265	-51.505	1.00	32.98	T1
	ATOM	369	CG	LYS	A	47	-16.284	80.557	-51.891	1.00	30.28	T1
	ATOM	370	CD	LYS	A	47	-16.233	80.143	-53.348	1.00	31.06	T1
	ATOM	371	CE	LYS	A	47	-17.592	79.727	-53.877	1.00	30.30	T1
60	ATOM	372	NZ	LYS	A	47	-18.513	80.890	-54.096	1.00	26.77	T1
	ATOM	373	C	LYS	A	47	-12.555	81.443	-51.939	1.00	28.11	T1
	ATOM	374	O	LYS	A	47	-12.524	82.282	-52.834	1.00	22.73	T1
	ATOM	375	N	GLU	A	48	-11.597	81.330	-51.032	1.00	31.40	T1
	ATOM	376	CA	GLU	A	48	-10.420	82.193	-51.044	1.00	36.50	T1
65	ATOM	377	CB	GLU	A	48	-10.577	83.348	-50.069	1.00	25.85	T1
	ATOM	378	CG	GLU	A	48	-11.581	84.386	-50.464	1.00	31.43	T1

	ATOM	379	CD	GLU	A	48	-11.774	85.422	-49.366	1.00	28.57	T1
	ATOM	380	OE1	GLU	A	48	-10.755	85.835	-48.750	1.00	32.19	T1
	ATOM	381	OE2	GLU	A	48	-12.941	85.822	-49.124	1.00	34.76	T1
	ATOM	382	C	GLU	A	48	-9.258	81.359	-50.571	1.00	30.52	T1
5	ATOM	383	O	GLU	A	48	-9.329	80.768	-49.491	1.00	29.50	T1
	ATOM	384	N	THR	A	49	-8.180	81.307	-51.347	1.00	26.70	T1
	ATOM	385	CA	THR	A	49	-7.047	80.503	-50.917	1.00	30.57	T1
	ATOM	386	CB	THR	A	49	-6.026	80.328	-52.037	1.00	32.94	T1
	ATOM	387	OG1	THR	A	49	-5.073	81.384	-51.964	1.00	34.31	T1
10	ATOM	388	CG2	THR	A	49	-6.712	80.368	-53.393	1.00	29.04	T1
	ATOM	389	C	THR	A	49	-6.394	81.177	-49.715	1.00	30.77	T1
	ATOM	390	O	THR	A	49	-6.440	82.395	-49.570	1.00	36.97	T1
	ATOM	391	N	GLY	A	50	-5.811	80.369	-48.841	1.00	33.41	T1
	ATOM	392	CA	GLY	A	50	-5.162	80.899	-47.657	1.00	29.39	T1
15	ATOM	393	C	GLY	A	50	-4.917	79.791	-46.650	1.00	28.42	T1
	ATOM	394	O	GLY	A	50	-4.994	78.608	-46.995	1.00	24.97	T1
	ATOM	395	N	TYR	A	51	-4.619	80.165	-45.408	1.00	33.80	T1
	ATOM	396	CA	TYR	A	51	-4.370	79.188	-44.347	1.00	32.60	T1
	ATOM	397	CB	TYR	A	51	-3.148	79.596	-43.514	1.00	26.81	T1
20	ATOM	398	CG	TYR	A	51	-1.853	79.350	-44.247	1.00	32.85	T1
	ATOM	399	CD1	TYR	A	51	-1.401	80.232	-45.226	1.00	34.53	T1
	ATOM	400	CE1	TYR	A	51	-0.267	79.944	-45.989	1.00	28.14	T1
	ATOM	401	CD2	TYR	A	51	-1.133	78.175	-44.037	1.00	29.75	T1
	ATOM	402	CE2	TYR	A	51	-0.005	77.874	-44.788	1.00	25.72	T1
25	ATOM	403	CZ	TYR	A	51	0.424	78.757	-45.769	1.00	33.16	T1
	ATOM	404	OH	TYR	A	51	1.512	78.424	-46.563	1.00	30.37	T1
	ATOM	405	C	TYR	A	51	-5.589	79.049	-43.455	1.00	34.80	T1
	ATOM	406	O	TYR	A	51	-6.111	80.035	-42.947	1.00	23.33	T1
	ATOM	407	N	PHE	A	52	-6.042	77.817	-43.265	1.00	28.64	T1
30	ATOM	408	CA	PHE	A	52	-7.216	77.581	-42.446	1.00	32.22	T1
	ATOM	409	CB	PHE	A	52	-8.363	77.038	-43.303	1.00	27.68	T1
	ATOM	410	CG	PHE	A	52	-8.758	77.929	-44.442	1.00	37.35	T1
	ATOM	411	CD1	PHE	A	52	-7.972	78.010	-45.583	1.00	31.89	T1
	ATOM	412	CD2	PHE	A	52	-9.932	78.674	-44.381	1.00	32.37	T1
35	ATOM	413	CE1	PHE	A	52	-8.351	78.820	-46.645	1.00	25.95	T1
	ATOM	414	CE2	PHE	A	52	-10.316	79.484	-45.436	1.00	36.15	T1
	ATOM	415	CZ	PHE	A	52	-9.525	79.558	-46.571	1.00	32.55	T1
	ATOM	416	C	PHE	A	52	-6.994	76.613	-41.291	1.00	29.88	T1
	ATOM	417	O	PHE	A	52	-6.190	75.686	-41.381	1.00	25.92	T1
40	ATOM	418	N	PHE	A	53	-7.720	76.852	-40.203	1.00	33.00	T1
	ATOM	419	CA	PHE	A	53	-7.690	75.992	-39.032	1.00	36.23	T1
	ATOM	420	CB	PHE	A	53	-7.859	76.807	-37.765	1.00	31.76	T1
	ATOM	421	CG	PHE	A	53	-8.056	75.973	-36.535	1.00	30.10	T1
	ATOM	422	CD1	PHE	A	53	-7.029	75.182	-36.048	1.00	29.49	T1
45	ATOM	423	CD2	PHE	A	53	-9.275	75.973	-35.864	1.00	32.85	T1
	ATOM	424	CE1	PHE	A	53	-7.216	74.400	-34.904	1.00	33.14	T1
	ATOM	425	CE2	PHE	A	53	-9.468	75.196	-34.724	1.00	39.17	T1
	ATOM	426	CZ	PHE	A	53	-8.442	74.411	-34.245	1.00	32.59	T1
	ATOM	427	C	PHE	A	53	-8.925	75.138	-39.241	1.00	35.45	T1
50	ATOM	428	O	PHE	A	53	-10.018	75.671	-39.387	1.00	29.46	T1
	ATOM	429	N	ILE	A	54	-8.760	73.823	-39.263	1.00	39.28	T1
	ATOM	430	CA	ILE	A	54	-9.889	72.937	-39.501	1.00	30.00	T1
	ATOM	431	CB	ILE	A	54	-9.665	72.149	-40.796	1.00	24.87	T1
	ATOM	432	CG2	ILE	A	54	-10.917	71.378	-41.167	1.00	36.42	T1
55	ATOM	433	CG1	ILE	A	54	-9.290	73.120	-41.910	1.00	32.77	T1
	ATOM	434	CD1	ILE	A	54	-8.617	72.470	-43.066	1.00	31.06	T1
	ATOM	435	C	ILE	A	54	-10.086	71.977	-38.343	1.00	31.23	T1
	ATOM	436	O	ILE	A	54	-9.134	71.338	-37.896	1.00	35.68	T1
	ATOM	437	N	TYR	A	55	-11.323	71.875	-37.863	1.00	29.04	T1
60	ATOM	438	CA	TYR	A	55	-11.620	70.995	-36.745	1.00	33.48	T1
	ATOM	439	CB	TYR	A	55	-11.833	71.822	-35.483	1.00	25.20	T1
	ATOM	440	CG	TYR	A	55	-12.915	72.858	-35.609	1.00	30.90	T1
	ATOM	441	CD1	TYR	A	55	-14.205	72.598	-35.165	1.00	29.43	T1
	ATOM	442	CE1	TYR	A	55	-15.212	73.543	-35.301	1.00	29.40	T1
65	ATOM	443	CD2	TYR	A	55	-12.652	74.093	-36.195	1.00	31.38	T1
	ATOM	444	CE2	TYR	A	55	-13.648	75.046	-36.339	1.00	22.87	T1

	ATOM	445	CZ	TYR	A	55	-14.927	74.767	-35.893	1.00	30.01	T1
	ATOM	446	OH	TYR	A	55	-15.916	75.710	-36.065	1.00	34.01	T1
	ATOM	447	C	TYR	A	55	-12.825	70.119	-36.995	1.00	25.58	T1
	ATOM	448	O	TYR	A	55	-13.517	70.279	-37.980	1.00	33.26	T1
5	ATOM	449	N	GLY	A	56	-13.063	69.174	-36.099	1.00	32.62	T1
	ATOM	450	CA	GLY	A	56	-14.196	68.287	-36.260	1.00	34.90	T1
	ATOM	451	C	GLY	A	56	-14.262	67.248	-35.159	1.00	30.65	T1
	ATOM	452	O	GLY	A	56	-13.236	66.718	-34.741	1.00	26.45	T1
	ATOM	453	N	GLN	A	57	-15.472	66.975	-34.681	1.00	31.77	T1
10	ATOM	454	CA	GLN	A	57	-15.694	65.982	-33.637	1.00	29.60	T1
	ATOM	455	CB	GLN	A	57	-15.974	66.655	-32.291	1.00	23.36	T1
	ATOM	456	CG	GLN	A	57	-16.377	65.670	-31.182	1.00	38.31	T1
	ATOM	457	CD	GLN	A	57	-16.520	66.314	-29.808	1.00	34.93	T1
	ATOM	458	OE1	GLN	A	57	-15.538	66.717	-29.187	1.00	33.10	T1
15	ATOM	459	NE2	GLN	A	57	-17.749	66.406	-29.330	1.00	31.14	T1
	ATOM	460	C	GLN	A	57	-16.875	65.101	-33.998	1.00	29.91	T1
	ATOM	461	O	GLN	A	57	-17.821	65.555	-34.639	1.00	26.30	T1
	ATOM	462	N	VAL	A	58	-16.807	63.841	-33.580	1.00	31.87	T1
	ATOM	463	CA	VAL	A	58	-17.857	62.860	-33.833	1.00	31.54	T1
20	ATOM	464	CB	VAL	A	58	-17.482	61.923	-35.004	1.00	32.11	T1
	ATOM	465	CG1	VAL	A	58	-18.450	60.757	-35.067	1.00	26.56	T1
	ATOM	466	CG2	VAL	A	58	-17.505	62.688	-36.313	1.00	26.98	T1
	ATOM	467	C	VAL	A	58	-18.036	62.007	-32.590	1.00	24.89	T1
	ATOM	468	O	VAL	A	58	-17.061	61.658	-31.946	1.00	28.22	T1
25	ATOM	469	N	LEU	A	59	-19.278	61.679	-32.247	1.00	37.54	T1
	ATOM	470	CA	LEU	A	59	-19.546	60.838	-31.080	1.00	32.69	T1
	ATOM	471	CB	LEU	A	59	-20.761	61.351	-30.303	1.00	38.29	T1
	ATOM	472	CG	LEU	A	59	-20.849	60.965	-28.823	1.00	30.05	T1
	ATOM	473	CD1	LEU	A	59	-22.277	61.167	-28.344	1.00	26.42	T1
30	ATOM	474	CD2	LEU	A	59	-20.441	59.520	-28.611	1.00	31.77	T1
	ATOM	475	C	LEU	A	59	-19.830	59.418	-31.576	1.00	30.26	T1
	ATOM	476	O	LEU	A	59	-20.839	59.174	-32.252	1.00	31.78	T1
	ATOM	477	N	TYR	A	60	-18.941	58.484	-31.246	1.00	26.50	T1
	ATOM	478	CA	TYR	A	60	-19.119	57.099	-31.671	1.00	26.97	T1
35	ATOM	479	CB	TYR	A	60	-17.770	56.462	-31.952	1.00	32.65	T1
	ATOM	480	CG	TYR	A	60	-17.033	57.194	-33.023	1.00	27.21	T1
	ATOM	481	CD1	TYR	A	60	-15.949	58.012	-32.715	1.00	32.97	T1
	ATOM	482	CE1	TYR	A	60	-15.304	58.747	-33.702	1.00	29.33	T1
	ATOM	483	CD2	TYR	A	60	-17.454	57.123	-34.341	1.00	23.99	T1
40	ATOM	484	CE2	TYR	A	60	-16.828	57.847	-35.332	1.00	34.87	T1
	ATOM	485	CZ	TYR	A	60	-15.755	58.659	-35.012	1.00	24.89	T1
	ATOM	486	OH	TYR	A	60	-15.155	59.395	-36.011	1.00	31.30	T1
	ATOM	487	C	TYR	A	60	-19.880	56.229	-30.677	1.00	28.77	T1
	ATOM	488	O	TYR	A	60	-19.516	56.128	-29.502	1.00	26.17	T1
45	ATOM	489	N	THR	A	61	-20.938	55.593	-31.162	1.00	28.11	T1
	ATOM	490	CA	THR	A	61	-21.742	54.720	-30.321	1.00	29.15	T1
	ATOM	491	CB	THR	A	61	-23.188	55.221	-30.235	1.00	31.21	T1
	ATOM	492	OG1	THR	A	61	-23.718	55.388	-31.560	1.00	31.05	T1
	ATOM	493	CG2	THR	A	61	-23.238	56.547	-29.492	1.00	38.00	T1
50	ATOM	494	C	THR	A	61	-21.714	53.326	-30.922	1.00	31.08	T1
	ATOM	495	O	THR	A	61	-22.520	52.469	-30.587	1.00	33.72	T1
	ATOM	496	N	ASP	A	62	-20.763	53.122	-31.822	1.00	31.05	T1
	ATOM	497	CA	ASP	A	62	-20.565	51.846	-32.488	1.00	34.03	T1
	ATOM	498	CB	ASP	A	62	-19.894	52.102	-33.836	1.00	30.78	T1
55	ATOM	499	CG	ASP	A	62	-19.929	50.899	-34.744	1.00	31.68	T1
	ATOM	500	OD1	ASP	A	62	-20.345	51.068	-35.925	1.00	32.63	T1
	ATOM	501	OD2	ASP	A	62	-19.537	49.800	-34.276	1.00	30.56	T1
	ATOM	502	C	ASP	A	62	-19.656	51.010	-31.579	1.00	29.75	T1
	ATOM	503	O	ASP	A	62	-18.813	51.561	-30.861	1.00	29.83	T1
60	ATOM	504	N	LYS	A	63	-19.813	49.692	-31.593	1.00	23.03	T1
	ATOM	505	CA	LYS	A	63	-18.968	48.871	-30.727	1.00	30.79	T1
	ATOM	506	CB	LYS	A	63	-19.835	47.921	-29.895	1.00	24.36	T1
	ATOM	507	CG	LYS	A	63	-20.672	46.953	-30.728	1.00	29.70	T1
	ATOM	508	CD	LYS	A	63	-21.486	46.009	-29.832	1.00	31.98	T1
65	ATOM	509	CE	LYS	A	63	-22.459	46.782	-28.923	1.00	35.27	T1
	ATOM	510	NZ	LYS	A	63	-23.243	45.884	-28.022	1.00	36.69	T1

	ATOM	511	C	LYS	A	63	-17.899	48.061	-31.464	1.00	26.50	T1
	ATOM	512	O	LYS	A	63	-17.392	47.061	-30.937	1.00	25.21	T1
	ATOM	513	N	THR	A	64	-17.528	48.491	-32.662	1.00	36.71	T1
	ATOM	514	CA	THR	A	64	-16.537	47.728	-33.399	1.00	36.05	T1
5	ATOM	515	CB	THR	A	64	-16.756	47.841	-34.936	1.00	32.42	T1
	ATOM	516	OG1	THR	A	64	-16.855	49.216	-35.322	1.00	26.37	T1
	ATOM	517	CG2	THR	A	64	-18.029	47.099	-35.338	1.00	40.39	T1
	ATOM	518	C	THR	A	64	-15.069	48.002	-33.067	1.00	31.32	T1
	ATOM	519	O	THR	A	64	-14.291	48.404	-33.931	1.00	32.64	T1
10	ATOM	520	N	TYR	A	65	-14.705	47.766	-31.809	1.00	35.20	T1
	ATOM	521	CA	TYR	A	65	-13.332	47.916	-31.313	1.00	31.02	T1
	ATOM	522	CB	TYR	A	65	-12.518	46.670	-31.694	1.00	35.17	T1
	ATOM	523	CG	TYR	A	65	-11.578	46.863	-32.868	1.00	32.56	T1
	ATOM	524	CD1	TYR	A	65	-10.244	47.243	-32.671	1.00	29.80	T1
15	ATOM	525	CE1	TYR	A	65	-9.376	47.423	-33.761	1.00	25.93	T1
	ATOM	526	CD2	TYR	A	65	-12.026	46.669	-34.182	1.00	25.36	T1
	ATOM	527	CE2	TYR	A	65	-11.172	46.848	-35.281	1.00	28.88	T1
	ATOM	528	CZ	TYR	A	65	-9.853	47.224	-35.069	1.00	22.04	T1
	ATOM	529	OH	TYR	A	65	-9.031	47.410	-36.168	1.00	28.62	T1
20	ATOM	530	C	TYR	A	65	-12.519	49.165	-31.671	1.00	39.45	T1
	ATOM	531	O	TYR	A	65	-11.576	49.514	-30.951	1.00	30.06	T1
	ATOM	532	N	ALA	A	66	-12.850	49.828	-32.773	1.00	30.09	T1
	ATOM	533	CA	ALA	A	66	-12.105	51.018	-33.163	1.00	33.00	T1
	ATOM	534	CB	ALA	A	66	-10.762	50.616	-33.743	1.00	27.69	T1
25	ATOM	535	C	ALA	A	66	-12.869	51.872	-34.164	1.00	30.55	T1
	ATOM	536	O	ALA	A	66	-13.169	51.425	-35.276	1.00	27.92	T1
	ATOM	537	N	MET	A	67	-13.188	53.099	-33.763	1.00	32.48	T1
	ATOM	538	CA	MET	A	67	-13.902	54.012	-34.644	1.00	28.97	T1
	ATOM	539	CB	MET	A	67	-15.250	54.393	-34.033	1.00	37.47	T1
30	ATOM	540	CG	MET	A	67	-16.254	53.254	-34.006	1.00	34.02	T1
	ATOM	541	SD	MET	A	67	-16.629	52.617	-35.669	1.00	34.29	T1
	ATOM	542	CE	MET	A	67	-17.780	53.858	-36.266	1.00	30.81	T1
	ATOM	543	C	MET	A	67	-13.059	55.257	-34.862	1.00	29.24	T1
	ATOM	544	O	MET	A	67	-12.062	55.457	-34.165	1.00	27.83	T1
35	ATOM	545	N	GLY	A	68	-13.450	56.085	-35.824	1.00	32.37	T1
	ATOM	546	CA	GLY	A	68	-12.703	57.302	-36.087	1.00	28.55	T1
	ATOM	547	C	GLY	A	68	-13.139	57.976	-37.367	1.00	31.40	T1
	ATOM	548	O	GLY	A	68	-13.916	57.407	-38.134	1.00	27.36	T1
	ATOM	549	N	HIS	A	69	-12.668	59.196	-37.601	1.00	33.39	T1
40	ATOM	550	CA	HIS	A	69	-13.027	59.908	-38.825	1.00	28.87	T1
	ATOM	551	CB	HIS	A	69	-14.142	60.938	-38.572	1.00	24.49	T1
	ATOM	552	CG	HIS	A	69	-13.849	61.909	-37.472	1.00	31.76	T1
	ATOM	553	CD2	HIS	A	69	-13.482	63.209	-37.506	1.00	31.88	T1
	ATOM	554	ND1	HIS	A	69	-13.954	61.577	-36.142	1.00	25.43	T1
45	ATOM	555	CE1	HIS	A	69	-13.668	62.632	-35.401	1.00	25.49	T1
	ATOM	556	NE2	HIS	A	69	-13.378	63.635	-36.206	1.00	29.23	T1
	ATOM	557	C	HIS	A	69	-11.833	60.587	-39.482	1.00	30.01	T1
	ATOM	558	O	HIS	A	69	-10.757	60.688	-38.896	1.00	30.93	T1
	ATOM	559	N	LEU	A	70	-12.039	61.046	-40.709	1.00	28.65	T1
50	ATOM	560	CA	LEU	A	70	-10.997	61.703	-41.477	1.00	27.23	T1
	ATOM	561	CB	LEU	A	70	-10.664	60.885	-42.724	1.00	29.19	T1
	ATOM	562	CG	LEU	A	70	-10.605	59.367	-42.613	1.00	33.10	T1
	ATOM	563	CD1	LEU	A	70	-10.473	58.788	-43.997	1.00	27.68	T1
	ATOM	564	CD2	LEU	A	70	-9.446	58.950	-41.751	1.00	24.67	T1
55	ATOM	565	C	LEU	A	70	-11.466	63.060	-41.949	1.00	27.27	T1
	ATOM	566	O	LEU	A	70	-12.636	63.238	-42.270	1.00	31.63	T1
	ATOM	567	N	ILE	A	71	-10.552	64.019	-41.992	1.00	31.95	T1
	ATOM	568	CA	ILE	A	71	-10.880	65.339	-42.504	1.00	28.20	T1
	ATOM	569	CB	ILE	A	71	-10.421	66.449	-41.553	1.00	30.55	T1
60	ATOM	570	CG2	ILE	A	71	-10.486	67.786	-42.245	1.00	29.76	T1
	ATOM	571	CG1	ILE	A	71	-11.323	66.472	-40.321	1.00	21.23	T1
	ATOM	572	CD1	ILE	A	71	-10.918	67.506	-39.287	1.00	24.23	T1
	ATOM	573	C	ILE	A	71	-10.102	65.386	-43.810	1.00	31.62	T1
	ATOM	574	O	ILE	A	71	-8.873	65.442	-43.813	1.00	30.07	T1
65	ATOM	575	N	GLN	A	72	-10.816	65.350	-44.928	1.00	35.77	T1
	ATOM	576	CA	GLN	A	72	-10.147	65.330	-46.220	1.00	32.31	T1

	ATOM	577	CB	GLN	A	72	-10.634	64.127	-47.008	1.00	26.47	T1
	ATOM	578	CG	GLN	A	72	-10.671	62.872	-46.161	1.00	30.39	T1
	ATOM	579	CD	GLN	A	72	-10.960	61.648	-46.981	1.00	23.69	T1
	ATOM	580	OE1	GLN	A	72	-11.961	61.592	-47.704	1.00	28.88	T1
5	ATOM	581	NE2	GLN	A	72	-10.084	60.652	-46.883	1.00	32.06	T1
	ATOM	582	C	GLN	A	72	-10.260	66.571	-47.075	1.00	28.96	T1
	ATOM	583	O	GLN	A	72	-11.182	67.371	-46.925	1.00	30.73	T1
	ATOM	584	N	ARG	A	73	-9.304	66.704	-47.986	1.00	31.30	T1
	ATOM	585	CA	ARG	A	73	-9.235	67.836	-48.886	1.00	40.24	T1
10	ATOM	586	CB	ARG	A	73	-7.953	68.607	-48.607	1.00	36.60	T1
	ATOM	587	CG	ARG	A	73	-7.765	69.794	-49.497	1.00	32.13	T1
	ATOM	588	CD	ARG	A	73	-6.312	70.154	-49.617	1.00	27.89	T1
	ATOM	589	NE	ARG	A	73	-6.115	71.298	-50.495	1.00	29.76	T1
	ATOM	590	CZ	ARG	A	73	-4.940	71.653	-50.994	1.00	27.66	T1
15	ATOM	591	NH1	ARG	A	73	-3.855	70.950	-50.706	1.00	26.51	T1
	ATOM	592	NH2	ARG	A	73	-4.850	72.717	-51.775	1.00	28.61	T1
	ATOM	593	C	ARG	A	73	-9.249	67.380	-50.348	1.00	28.16	T1
	ATOM	594	O	ARG	A	73	-8.523	66.451	-50.715	1.00	23.44	T1
	ATOM	595	N	LYS	A	74	-10.077	68.019	-51.175	1.00	34.41	T1
20	ATOM	596	CA	LYS	A	74	-10.152	67.695	-52.598	1.00	29.07	T1
	ATOM	597	CB	LYS	A	74	-11.599	67.571	-53.063	1.00	34.35	T1
	ATOM	598	CG	LYS	A	74	-12.336	66.374	-52.497	1.00	30.85	T1
	ATOM	599	CD	LYS	A	74	-13.821	66.348	-52.922	1.00	28.70	T1
	ATOM	600	CE	LYS	A	74	-14.563	65.136	-52.318	1.00	28.03	T1
25	ATOM	601	NZ	LYS	A	74	-15.999	65.039	-52.744	1.00	34.06	T1
	ATOM	602	C	LYS	A	74	-9.492	68.819	-53.374	1.00	27.44	T1
	ATOM	603	O	LYS	A	74	-10.100	69.871	-53.589	1.00	31.06	T1
	ATOM	604	N	LYS	A	75	-8.251	68.600	-53.792	1.00	26.43	T1
	ATOM	605	CA	LYS	A	75	-7.497	69.604	-54.540	1.00	26.50	T1
30	ATOM	606	CB	LYS	A	75	-6.096	69.081	-54.877	1.00	30.89	T1
	ATOM	607	CG	LYS	A	75	-5.218	68.727	-53.697	1.00	37.11	T1
	ATOM	608	CD	LYS	A	75	-3.872	68.226	-54.190	1.00	40.43	T1
	ATOM	609	CE	LYS	A	75	-2.966	67.836	-53.028	1.00	31.87	T1
	ATOM	610	NZ	LYS	A	75	-1.646	67.231	-53.438	1.00	28.85	T1
35	ATOM	611	C	LYS	A	75	-8.192	69.978	-55.843	1.00	34.46	T1
	ATOM	612	O	LYS	A	75	-8.662	69.101	-56.563	1.00	27.37	T1
	ATOM	613	N	VAL	A	76	-8.242	71.272	-56.152	1.00	36.10	T1
	ATOM	614	CA	VAL	A	76	-8.858	71.730	-57.398	1.00	34.84	T1
	ATOM	615	CB	VAL	A	76	-9.125	73.228	-57.413	1.00	33.91	T1
40	ATOM	616	CG1	VAL	A	76	-10.103	73.549	-58.498	1.00	32.01	T1
	ATOM	617	CG2	VAL	A	76	-9.617	73.683	-56.093	1.00	31.96	T1
	ATOM	618	C	VAL	A	76	-7.845	71.496	-58.501	1.00	29.58	T1
	ATOM	619	O	VAL	A	76	-8.186	71.075	-59.607	1.00	31.58	T1
	ATOM	620	N	HIS	A	77	-6.593	71.798	-58.183	1.00	29.25	T1
45	ATOM	621	CA	HIS	A	77	-5.504	71.635	-59.120	1.00	30.42	T1
	ATOM	622	CB	HIS	A	77	-4.628	72.878	-59.088	1.00	28.32	T1
	ATOM	623	CG	HIS	A	77	-5.371	74.136	-59.407	1.00	25.36	T1
	ATOM	624	CD2	HIS	A	77	-5.079	75.434	-59.153	1.00	35.54	T1
	ATOM	625	ND1	HIS	A	77	-6.539	74.142	-60.144	1.00	28.93	T1
50	ATOM	626	CE1	HIS	A	77	-6.931	75.388	-60.338	1.00	31.96	T1
	ATOM	627	NE2	HIS	A	77	-6.062	76.191	-59.747	1.00	29.26	T1
	ATOM	628	C	HIS	A	77	-4.707	70.397	-58.750	1.00	26.67	T1
	ATOM	629	O	HIS	A	77	-4.579	70.072	-57.570	1.00	29.56	T1
	ATOM	630	N	VAL	A	78	-4.154	69.712	-59.748	1.00	27.67	T1
55	ATOM	631	CA	VAL	A	78	-3.425	68.494	-59.450	1.00	25.24	T1
	ATOM	632	CB	VAL	A	78	-4.244	67.282	-59.931	1.00	32.89	T1
	ATOM	633	CG1	VAL	A	78	-3.590	65.994	-59.499	1.00	28.47	T1
	ATOM	634	CG2	VAL	A	78	-5.639	67.347	-59.322	1.00	30.58	T1
	ATOM	635	C	VAL	A	78	-1.952	68.346	-59.875	1.00	27.67	T1
60	ATOM	636	O	VAL	A	78	-1.077	68.281	-59.000	1.00	35.37	T1
	ATOM	637	N	PHE	A	79	-1.644	68.285	-61.168	1.00	33.36	T1
	ATOM	638	CA	PHE	A	79	-0.232	68.119	-61.588	1.00	23.78	T1
	ATOM	639	CB	PHE	A	79	0.733	69.006	-60.780	1.00	28.94	T1
	ATOM	640	CG	PHE	A	79	0.361	70.450	-60.746	1.00	30.56	T1
65	ATOM	641	CD1	PHE	A	79	-0.275	70.989	-59.631	1.00	27.92	T1
	ATOM	642	CD2	PHE	A	79	0.636	71.272	-61.829	1.00	28.00	T1

	ATOM	643	CE1	PHE	A	79	-0.635	72.323	-59.594	1.00	31.49	T1
	ATOM	644	CE2	PHE	A	79	0.280	72.612	-61.807	1.00	36.51	T1
	ATOM	645	CZ	PHE	A	79	-0.358	73.139	-60.684	1.00	25.74	T1
	ATOM	646	C	PHE	A	79	0.312	66.688	-61.465	1.00	31.62	T1
5	ATOM	647	O	PHE	A	79	0.314	66.096	-60.385	1.00	29.63	T1
	ATOM	648	N	GLY	A	80	0.803	66.158	-62.580	1.00	26.96	T1
	ATOM	649	CA	GLY	A	80	1.387	64.826	-62.604	1.00	23.46	T1
	ATOM	650	C	GLY	A	80	0.610	63.725	-61.919	1.00	28.46	T1
	ATOM	651	O	GLY	A	80	-0.588	63.561	-62.151	1.00	29.44	T1
10	ATOM	652	N	ASP	A	81	1.307	62.971	-61.070	1.00	26.00	T1
	ATOM	653	CA	ASP	A	81	0.714	61.856	-60.343	1.00	32.30	T1
	ATOM	654	CB	ASP	A	81	1.689	60.675	-60.331	1.00	27.93	T1
	ATOM	655	CG	ASP	A	81	2.941	60.956	-59.531	1.00	40.18	T1
	ATOM	656	OD1	ASP	A	81	3.243	62.132	-59.265	1.00	28.59	T1
15	ATOM	657	OD2	ASP	A	81	3.640	59.993	-59.172	1.00	27.04	T1
	ATOM	658	C	ASP	A	81	0.266	62.183	-58.912	1.00	29.96	T1
	ATOM	659	O	ASP	A	81	0.278	61.314	-58.033	1.00	21.26	T1
	ATOM	660	N	GLU	A	82	-0.123	63.432	-58.672	1.00	28.53	T1
	ATOM	661	CA	GLU	A	82	-0.593	63.811	-57.348	1.00	36.31	T1
20	ATOM	662	CB	GLU	A	82	-0.801	65.315	-57.241	1.00	29.69	T1
	ATOM	663	CG	GLU	A	82	0.404	66.154	-57.015	1.00	34.40	T1
	ATOM	664	CD	GLU	A	82	0.025	67.433	-56.285	1.00	30.81	T1
	ATOM	665	OE1	GLU	A	82	-1.001	68.043	-56.654	1.00	34.76	T1
	ATOM	666	OE2	GLU	A	82	0.741	67.837	-55.337	1.00	27.76	T1
25	ATOM	667	C	GLU	A	82	-1.958	63.169	-57.163	1.00	30.42	T1
	ATOM	668	O	GLU	A	82	-2.675	62.963	-58.143	1.00	31.60	T1
	ATOM	669	N	LEU	A	83	-2.320	62.844	-55.927	1.00	31.31	T1
	ATOM	670	CA	LEU	A	83	-3.647	62.301	-55.678	1.00	37.43	T1
	ATOM	671	CB	LEU	A	83	-3.660	61.343	-54.489	1.00	23.70	T1
30	ATOM	672	CG	LEU	A	83	-2.950	60.004	-54.665	1.00	36.21	T1
	ATOM	673	CD1	LEU	A	83	-1.462	60.230	-54.877	1.00	26.80	T1
	ATOM	674	CD2	LEU	A	83	-3.182	59.155	-53.434	1.00	36.10	T1
	ATOM	675	C	LEU	A	83	-4.416	63.555	-55.333	1.00	34.44	T1
	ATOM	676	O	LEU	A	83	-3.958	64.353	-54.520	1.00	24.85	T1
35	ATOM	677	N	SER	A	84	-5.569	63.750	-55.959	1.00	26.06	T1
	ATOM	678	CA	SER	A	84	-6.354	64.952	-55.705	1.00	26.08	T1
	ATOM	679	CB	SER	A	84	-7.304	65.220	-56.873	1.00	37.72	T1
	ATOM	680	OG	SER	A	84	-7.993	64.044	-57.237	1.00	28.65	T1
	ATOM	681	C	SER	A	84	-7.130	64.909	-54.397	1.00	34.42	T1
40	ATOM	682	O	SER	A	84	-7.781	65.884	-54.022	1.00	36.43	T1
	ATOM	683	N	LEU	A	85	-7.057	63.781	-53.698	1.00	29.74	T1
	ATOM	684	CA	LEU	A	85	-7.751	63.642	-52.429	1.00	27.05	T1
	ATOM	685	CB	LEU	A	85	-8.730	62.465	-52.494	1.00	31.59	T1
	ATOM	686	CG	LEU	A	85	-9.848	62.326	-51.445	1.00	25.64	T1
45	ATOM	687	CD1	LEU	A	85	-9.276	61.941	-50.088	1.00	27.95	T1
	ATOM	688	CD2	LEU	A	85	-10.633	63.628	-51.359	1.00	32.95	T1
	ATOM	689	C	LEU	A	85	-6.716	63.407	-51.345	1.00	34.31	T1
	ATOM	690	O	LEU	A	85	-6.121	62.335	-51.264	1.00	25.40	T1
	ATOM	691	N	VAL	A	86	-6.485	64.421	-50.520	1.00	22.34	T1
50	ATOM	692	CA	VAL	A	86	-5.514	64.307	-49.431	1.00	30.13	T1
	ATOM	693	CB	VAL	A	86	-4.558	65.516	-49.375	1.00	30.31	T1
	ATOM	694	CG1	VAL	A	86	-3.653	65.398	-48.163	1.00	32.35	T1
	ATOM	695	CG2	VAL	A	86	-3.739	65.596	-50.649	1.00	29.51	T1
	ATOM	696	C	VAL	A	86	-6.270	64.274	-48.126	1.00	32.65	T1
55	ATOM	697	O	VAL	A	86	-7.239	65.016	-47.944	1.00	30.66	T1
	ATOM	698	N	THR	A	87	-5.840	63.414	-47.215	1.00	26.19	T1
	ATOM	699	CA	THR	A	87	-6.507	63.356	-45.931	1.00	30.33	T1
	ATOM	700	CB	THR	A	87	-6.881	61.903	-45.546	1.00	33.67	T1
	ATOM	701	OG1	THR	A	87	-6.174	61.518	-44.366	1.00	39.32	T1
60	ATOM	702	CG2	THR	A	87	-6.556	60.948	-46.680	1.00	24.27	T1
	ATOM	703	C	THR	A	87	-5.599	63.984	-44.884	1.00	30.83	T1
	ATOM	704	O	THR	A	87	-4.482	63.532	-44.657	1.00	34.62	T1
	ATOM	705	N	LEU	A	88	-6.082	65.063	-44.285	1.00	39.24	T1
	ATOM	706	CA	LEU	A	88	-5.353	65.789	-43.253	1.00	38.69	T1
65	ATOM	707	CB	LEU	A	88	-5.658	67.285	-43.363	1.00	20.33	T1
	ATOM	708	CG	LEU	A	88	-5.471	68.135	-44.625	1.00	31.82	T1

	ATOM	709	CD1	LEU	A	88	-5.606	67.311	-45.857	1.00	30.40	T1
	ATOM	710	CD2	LEU	A	88	-6.502	69.240	-44.633	1.00	26.62	T1
	ATOM	711	C	LEU	A	88	-5.878	65.310	-41.898	1.00	30.10	T1
	ATOM	712	O	LEU	A	88	-7.064	64.972	-41.752	1.00	29.30	T1
5	ATOM	713	N	PHE	A	89	-5.024	65.243	-40.897	1.00	28.11	T1
	ATOM	714	CA	PHE	A	89	-5.532	64.872	-39.570	1.00	32.58	T1
	ATOM	715	CB	PHE	A	89	-6.599	65.893	-39.162	1.00	32.99	T1
	ATOM	716	CG	PHE	A	89	-6.210	67.300	-39.501	1.00	26.77	T1
	ATOM	717	CD1	PHE	A	89	-7.165	68.232	-39.901	1.00	24.25	T1
10	ATOM	718	CD2	PHE	A	89	-4.852	67.672	-39.517	1.00	28.47	T1
	ATOM	719	CE1	PHE	A	89	-6.774	69.514	-40.325	1.00	31.81	T1
	ATOM	720	CE2	PHE	A	89	-4.451	68.948	-39.937	1.00	24.54	T1
	ATOM	721	CZ	PHE	A	89	-5.413	69.867	-40.345	1.00	35.46	T1
	ATOM	722	C	PHE	A	89	-6.047	63.451	-39.314	1.00	34.60	T1
15	ATOM	723	O	PHE	A	89	-5.234	62.533	-39.170	1.00	34.92	T1
	ATOM	724	N	ARG	A	90	-7.364	63.251	-39.218	1.00	29.02	T1
	ATOM	725	CA	ARG	A	90	-7.869	61.897	-38.924	1.00	31.43	T1
	ATOM	726	CB	ARG	A	90	-7.303	60.903	-39.939	1.00	31.65	T1
	ATOM	727	CG	ARG	A	90	-7.151	59.487	-39.406	1.00	26.40	T1
20	ATOM	728	CD	ARG	A	90	-6.248	58.666	-40.297	1.00	30.25	T1
	ATOM	729	NE	ARG	A	90	-5.957	57.389	-39.667	1.00	33.59	T1
	ATOM	730	CZ	ARG	A	90	-4.785	56.767	-39.744	1.00	25.79	T1
	ATOM	731	NH1	ARG	A	90	-3.783	57.306	-40.434	1.00	31.49	T1
	ATOM	732	NH2	ARG	A	90	-4.611	55.609	-39.109	1.00	32.64	T1
25	ATOM	733	C	ARG	A	90	-7.532	61.373	-37.504	1.00	33.30	T1
	ATOM	734	O	ARG	A	90	-6.369	61.375	-37.090	1.00	34.26	T1
	ATOM	735	N	CYS	A	91	-8.545	60.894	-36.783	1.00	37.09	T1
	ATOM	736	CA	CYS	A	91	-8.351	60.360	-35.429	1.00	25.64	T1
	ATOM	737	CB	CYS	A	91	-8.921	61.329	-34.400	1.00	30.67	T1
30	ATOM	738	SG	CYS	A	91	-10.633	61.707	-34.709	1.00	27.00	T1
	ATOM	739	C	CYS	A	91	-8.993	58.971	-35.254	1.00	30.17	T1
	ATOM	740	O	CYS	A	91	-9.788	58.533	-36.085	1.00	30.65	T1
	ATOM	741	N	ILE	A	92	-8.640	58.287	-34.167	1.00	27.44	T1
	ATOM	742	CA	ILE	A	92	-9.139	56.937	-33.886	1.00	32.98	T1
35	ATOM	743	CB	ILE	A	92	-8.077	55.875	-34.221	1.00	28.20	T1
	ATOM	744	CG2	ILE	A	92	-8.628	54.489	-33.986	1.00	35.13	T1
	ATOM	745	CG1	ILE	A	92	-7.636	56.019	-35.673	1.00	36.46	T1
	ATOM	746	CD1	ILE	A	92	-6.371	55.238	-36.004	1.00	33.39	T1
	ATOM	747	C	ILE	A	92	-9.458	56.783	-32.405	1.00	28.29	T1
40	ATOM	748	O	ILE	A	92	-8.892	57.474	-31.569	1.00	29.55	T1
	ATOM	749	N	GLN	A	93	-10.358	55.867	-32.076	1.00	29.69	T1
	ATOM	750	CA	GLN	A	93	-10.724	55.629	-30.685	1.00	30.63	T1
	ATOM	751	CB	GLN	A	93	-11.839	56.586	-30.244	1.00	25.31	T1
	ATOM	752	CG	GLN	A	93	-11.356	57.926	-29.721	1.00	28.86	T1
45	ATOM	753	CD	GLN	A	93	-11.274	57.973	-28.202	1.00	32.46	T1
	ATOM	754	OE1	GLN	A	93	-10.365	57.401	-27.585	1.00	26.03	T1
	ATOM	755	NE2	GLN	A	93	-12.243	58.652	-27.587	1.00	35.40	T1
	ATOM	756	C	GLN	A	93	-11.193	54.191	-30.494	1.00	26.31	T1
	ATOM	757	O	GLN	A	93	-12.163	53.743	-31.138	1.00	26.62	T1
50	ATOM	758	N	ASN	A	94	-10.498	53.465	-29.621	1.00	35.07	T1
	ATOM	759	CA	ASN	A	94	-10.879	52.091	-29.335	1.00	26.14	T1
	ATOM	760	CB	ASN	A	94	-9.910	51.458	-28.327	1.00	30.46	T1
	ATOM	761	CG	ASN	A	94	-8.628	50.966	-28.977	1.00	27.64	T1
	ATOM	762	OD1	ASN	A	94	-8.671	50.200	-29.935	1.00	31.37	T1
55	ATOM	763	ND2	ASN	A	94	-7.487	51.395	-28.456	1.00	35.57	T1
	ATOM	764	C	ASN	A	94	-12.284	52.161	-28.747	1.00	33.11	T1
	ATOM	765	O	ASN	A	94	-12.626	53.125	-28.051	1.00	36.66	T1
	ATOM	766	N	MET	A	95	-13.102	51.156	-29.043	1.00	25.91	T1
	ATOM	767	CA	MET	A	95	-14.465	51.117	-28.532	1.00	26.46	T1
60	ATOM	768	CB	MET	A	95	-15.468	50.973	-29.684	1.00	27.82	T1
	ATOM	769	CG	MET	A	95	-15.446	52.120	-30.671	1.00	32.69	T1
	ATOM	770	SD	MET	A	95	-15.592	53.746	-29.844	1.00	23.80	T1
	ATOM	771	CE	MET	A	95	-17.366	53.800	-29.432	1.00	32.54	T1
	ATOM	772	C	MET	A	95	-14.649	49.952	-27.573	1.00	31.77	T1
65	ATOM	773	O	MET	A	95	-13.962	48.929	-27.680	1.00	25.24	T1
	ATOM	774	N	PRO	A	96	-15.577	50.101	-26.608	1.00	32.80	T1

	ATOM	775	CD	PRO	A	96	-16.337	51.319	-26.287	1.00	25.24	T1
	ATOM	776	CA	PRO	A	96	-15.865	49.051	-25.625	1.00	29.94	T1
	ATOM	777	CB	PRO	A	96	-16.666	49.774	-24.531	1.00	24.36	T1
	ATOM	778	CG	PRO	A	96	-16.401	51.231	-24.776	1.00	32.51	T1
5	ATOM	779	C	PRO	A	96	-16.753	48.061	-26.359	1.00	28.59	T1
	ATOM	780	O	PRO	A	96	-16.937	48.154	-27.584	1.00	29.89	T1
	ATOM	781	N	GLU	A	97	-17.333	47.134	-25.611	1.00	27.69	T1
	ATOM	782	CA	GLU	A	97	-18.195	46.147	-26.225	1.00	33.36	T1
	ATOM	783	CB	GLU	A	97	-17.687	44.755	-25.890	1.00	25.66	T1
10	ATOM	784	CG	GLU	A	97	-18.282	43.695	-26.769	1.00	28.10	T1
	ATOM	785	CD	GLU	A	97	-17.225	42.708	-27.224	1.00	27.26	T1
	ATOM	786	OE1	GLU	A	97	-16.610	42.061	-26.331	1.00	28.61	T1
	ATOM	787	OE2	GLU	A	97	-17.000	42.589	-28.467	1.00	28.33	T1
	ATOM	788	C	GLU	A	97	-19.618	46.316	-25.725	1.00	40.30	T1
15	ATOM	789	O	GLU	A	97	-20.576	45.916	-26.386	1.00	33.78	T1
	ATOM	790	N	THR	A	98	-19.743	46.940	-24.562	1.00	27.16	T1
	ATOM	791	CA	THR	A	98	-21.040	47.133	-23.950	1.00	30.67	T1
	ATOM	792	CB	THR	A	98	-20.932	47.001	-22.434	1.00	26.76	T1
	ATOM	793	OG1	THR	A	98	-20.075	48.045	-21.937	1.00	41.14	T1
20	ATOM	794	CG2	THR	A	98	-20.348	45.640	-22.066	1.00	29.00	T1
	ATOM	795	C	THR	A	98	-21.739	48.456	-24.260	1.00	31.97	T1
	ATOM	796	O	THR	A	98	-22.726	48.475	-25.008	1.00	30.46	T1
	ATOM	797	N	LEU	A	99	-21.251	49.562	-23.703	1.00	30.71	T1
	ATOM	798	CA	LEU	A	99	-21.936	50.821	-23.936	1.00	32.06	T1
25	ATOM	799	CB	LEU	A	99	-22.428	51.378	-22.602	1.00	29.91	T1
	ATOM	800	CG	LEU	A	99	-23.436	50.441	-21.921	1.00	33.46	T1
	ATOM	801	CD1	LEU	A	99	-23.783	50.937	-20.521	1.00	32.36	T1
	ATOM	802	CD2	LEU	A	99	-24.682	50.355	-22.792	1.00	32.97	T1
	ATOM	803	C	LEU	A	99	-21.126	51.861	-24.678	1.00	35.15	T1
30	ATOM	804	O	LEU	A	99	-20.710	52.872	-24.095	1.00	28.55	T1
	ATOM	805	N	PRO	A	100	-20.914	51.640	-25.989	1.00	31.25	T1
	ATOM	806	CD	PRO	A	100	-21.473	50.530	-26.782	1.00	32.35	T1
	ATOM	807	CA	PRO	A	100	-20.149	52.547	-26.852	1.00	28.32	T1
	ATOM	808	CB	PRO	A	100	-20.488	52.047	-28.257	1.00	26.69	T1
35	ATOM	809	CG	PRO	A	100	-20.623	50.576	-28.043	1.00	28.01	T1
	ATOM	810	C	PRO	A	100	-20.518	54.017	-26.656	1.00	24.82	T1
	ATOM	811	O	PRO	A	100	-21.679	54.416	-26.813	1.00	32.09	T1
	ATOM	812	N	ASN	A	101	-19.512	54.813	-26.315	1.00	30.61	T1
	ATOM	813	CA	ASN	A	101	-19.691	56.243	-26.094	1.00	25.74	T1
40	ATOM	814	CB	ASN	A	101	-20.342	56.495	-24.747	1.00	29.18	T1
	ATOM	815	CG	ASN	A	101	-21.818	56.466	-24.826	1.00	28.59	T1
	ATOM	816	OD1	ASN	A	101	-22.430	57.359	-25.418	1.00	30.91	T1
	ATOM	817	ND2	ASN	A	101	-22.422	55.429	-24.246	1.00	30.08	T1
	ATOM	818	C	ASN	A	101	-18.346	56.929	-26.090	1.00	26.46	T1
45	ATOM	819	O	ASN	A	101	-17.795	57.183	-25.007	1.00	29.86	T1
	ATOM	820	N	ASN	A	102	-17.802	57.246	-27.263	1.00	35.21	T1
	ATOM	821	CA	ASN	A	102	-16.511	57.884	-27.222	1.00	35.45	T1
	ATOM	822	CB	ASN	A	102	-15.447	56.952	-27.792	1.00	30.10	T1
	ATOM	823	CG	ASN	A	102	-14.942	55.968	-26.740	1.00	26.40	T1
50	ATOM	824	OD1	ASN	A	102	-14.665	56.355	-25.594	1.00	22.72	T1
	ATOM	825	ND2	ASN	A	102	-14.826	54.696	-27.114	1.00	31.16	T1
	ATOM	826	C	ASN	A	102	-16.303	59.304	-27.704	1.00	29.40	T1
	ATOM	827	O	ASN	A	102	-15.669	60.082	-26.978	1.00	29.70	T1
	ATOM	828	N	SER	A	103	-16.801	59.699	-28.870	1.00	26.26	T1
55	ATOM	829	CA	SER	A	103	-16.550	61.099	-29.244	1.00	26.89	T1
	ATOM	830	CB	SER	A	103	-17.193	62.046	-28.213	1.00	34.54	T1
	ATOM	831	OG	SER	A	103	-16.282	63.045	-27.781	1.00	25.40	T1
	ATOM	832	C	SER	A	103	-15.040	61.394	-29.337	1.00	31.26	T1
	ATOM	833	O	SER	A	103	-14.254	61.104	-28.432	1.00	36.38	T1
60	ATOM	834	N	CYS	A	104	-14.640	61.991	-30.443	1.00	34.68	T1
	ATOM	835	CA	CYS	A	104	-13.239	62.283	-30.641	1.00	24.17	T1
	ATOM	836	CB	CYS	A	104	-12.590	61.114	-31.375	1.00	28.98	T1
	ATOM	837	SG	CYS	A	104	-10.788	61.142	-31.419	1.00	26.11	T1
	ATOM	838	C	CYS	A	104	-13.086	63.567	-31.441	1.00	35.35	T1
65	ATOM	839	O	CYS	A	104	-13.670	63.722	-32.511	1.00	33.20	T1
	ATOM	840	N	TYR	A	105	-12.299	64.488	-30.903	1.00	35.21	T1

	ATOM	841	CA	TYR	A	105	-12.049	65.771	-31.544	1.00	34.06	T1
	ATOM	842	CB	TYR	A	105	-12.203	66.893	-30.518	1.00	34.49	T1
	ATOM	843	CG	TYR	A	105	-11.866	68.267	-31.036	1.00	27.80	T1
	ATOM	844	CD1	TYR	A	105	-12.856	69.108	-31.530	1.00	36.29	T1
5	ATOM	845	CE1	TYR	A	105	-12.548	70.392	-31.993	1.00	30.72	T1
	ATOM	846	CD2	TYR	A	105	-10.549	68.737	-31.022	1.00	29.27	T1
	ATOM	847	CE2	TYR	A	105	-10.228	70.022	-31.491	1.00	32.17	T1
	ATOM	848	CZ	TYR	A	105	-11.234	70.841	-31.969	1.00	32.74	T1
	ATOM	849	OH	TYR	A	105	-10.936	72.109	-32.402	1.00	32.48	T1
10	ATOM	850	C	TYR	A	105	-10.634	65.792	-32.098	1.00	36.15	T1
	ATOM	851	O	TYR	A	105	-9.729	65.193	-31.535	1.00	25.91	T1
	ATOM	852	N	SER	A	106	-10.446	66.488	-33.206	1.00	25.51	T1
	ATOM	853	CA	SER	A	106	-9.130	66.598	-33.797	1.00	30.66	T1
	ATOM	854	CB	SER	A	106	-8.788	65.329	-34.592	1.00	28.43	T1
15	ATOM	855	OG	SER	A	106	-7.423	65.314	-34.980	1.00	28.52	T1
	ATOM	856	C	SER	A	106	-9.178	67.818	-34.698	1.00	27.96	T1
	ATOM	857	O	SER	A	106	-10.211	68.105	-35.297	1.00	32.30	T1
	ATOM	858	N	ALA	A	107	-8.067	68.542	-34.772	1.00	33.29	T1
	ATOM	859	CA	ALA	A	107	-7.992	69.736	-35.597	1.00	31.63	T1
20	ATOM	860	CB	ALA	A	107	-8.586	70.915	-34.844	1.00	31.43	T1
	ATOM	861	C	ALA	A	107	-6.554	70.041	-36.001	1.00	36.43	T1
	ATOM	862	O	ALA	A	107	-5.614	69.508	-35.416	1.00	29.27	T1
	ATOM	863	N	GLY	A	108	-6.394	70.901	-37.004	1.00	23.03	T1
	ATOM	864	CA	GLY	A	108	-5.070	71.271	-37.465	1.00	29.99	T1
25	ATOM	865	C	GLY	A	108	-5.131	72.414	-38.457	1.00	29.99	T1
	ATOM	866	O	GLY	A	108	-6.214	72.893	-38.787	1.00	27.48	T1
	ATOM	867	N	ILE	A	109	-3.973	72.853	-38.935	1.00	30.89	T1
	ATOM	868	CA	ILE	A	109	-3.912	73.943	-39.901	1.00	23.10	T1
	ATOM	869	CB	ILE	A	109	-2.935	75.041	-39.447	1.00	30.72	T1
30	ATOM	870	CG2	ILE	A	109	-2.864	76.136	-40.501	1.00	36.69	T1
	ATOM	871	CG1	ILE	A	109	-3.383	75.620	-38.107	1.00	26.68	T1
	ATOM	872	CD1	ILE	A	109	-2.365	76.536	-37.483	1.00	33.42	T1
	ATOM	873	C	ILE	A	109	-3.433	73.406	-41.243	1.00	32.68	T1
	ATOM	874	O	ILE	A	109	-2.562	72.546	-41.294	1.00	29.21	T1
35	ATOM	875	N	ALA	A	110	-3.999	73.917	-42.327	1.00	32.51	T1
	ATOM	876	CA	ALA	A	110	-3.619	73.470	-43.655	1.00	32.54	T1
	ATOM	877	CB	ALA	A	110	-4.484	72.301	-44.078	1.00	30.87	T1
	ATOM	878	C	ALA	A	110	-3.791	74.614	-44.631	1.00	23.75	T1
	ATOM	879	O	ALA	A	110	-4.570	75.534	-44.380	1.00	31.86	T1
40	ATOM	880	N	LYS	A	111	-3.058	74.576	-45.738	1.00	26.07	T1
	ATOM	881	CA	LYS	A	111	-3.188	75.629	-46.725	1.00	29.40	T1
	ATOM	882	CB	LYS	A	111	-1.853	75.951	-47.367	1.00	36.99	T1
	ATOM	883	CG	LYS	A	111	-1.969	77.079	-48.370	1.00	31.67	T1
	ATOM	884	CD	LYS	A	111	-0.618	77.440	-48.983	1.00	29.35	T1
45	ATOM	885	CE	LYS	A	111	-0.749	78.608	-49.967	1.00	30.29	T1
	ATOM	886	NZ	LYS	A	111	0.580	79.015	-50.553	1.00	22.06	T1
	ATOM	887	C	LYS	A	111	-4.145	75.119	-47.775	1.00	32.23	T1
	ATOM	888	O	LYS	A	111	-3.996	73.993	-48.242	1.00	29.03	T1
	ATOM	889	N	LEU	A	112	-5.127	75.941	-48.136	1.00	31.54	T1
50	ATOM	890	CA	LEU	A	112	-6.138	75.570	-49.127	1.00	30.21	T1
	ATOM	891	CB	LEU	A	112	-7.495	75.426	-48.442	1.00	31.45	T1
	ATOM	892	CG	LEU	A	112	-7.540	74.586	-47.172	1.00	29.91	T1
	ATOM	893	CD1	LEU	A	112	-8.858	74.810	-46.456	1.00	34.32	T1
	ATOM	894	CD2	LEU	A	112	-7.350	73.136	-47.529	1.00	27.03	T1
55	ATOM	895	C	LEU	A	112	-6.247	76.629	-50.227	1.00	27.12	T1
	ATOM	896	O	LEU	A	112	-5.808	77.768	-50.051	1.00	31.38	T1
	ATOM	897	N	GLU	A	113	-6.848	76.253	-51.353	1.00	35.07	T1
	ATOM	898	CA	GLU	A	113	-7.026	77.176	-52.472	1.00	28.61	T1
	ATOM	899	CB	GLU	A	113	-6.191	76.764	-53.671	1.00	23.55	T1
60	ATOM	900	CG	GLU	A	113	-4.822	76.229	-53.379	1.00	31.18	T1
	ATOM	901	CD	GLU	A	113	-3.933	76.297	-54.618	1.00	30.29	T1
	ATOM	902	OE1	GLU	A	113	-4.403	75.927	-55.736	1.00	30.23	T1
	ATOM	903	OE2	GLU	A	113	-2.759	76.726	-54.467	1.00	26.25	T1
	ATOM	904	C	GLU	A	113	-8.465	77.233	-52.963	1.00	28.20	T1
65	ATOM	905	O	GLU	A	113	-9.240	76.297	-52.760	1.00	31.90	T1
	ATOM	906	N	GLU	A	114	-8.803	78.331	-53.635	1.00	29.69	T1

	ATOM	907	CA	GLU	A	114	-10.133	78.510	-54.207	1.00	33.14	T1
	ATOM	908	CB	GLU	A	114	-10.124	79.575	-55.283	1.00	32.10	T1
	ATOM	909	CG	GLU	A	114	-10.355	80.973	-54.830	1.00	28.54	T1
	ATOM	910	CD	GLU	A	114	-10.767	81.844	-56.001	1.00	34.74	T1
5	ATOM	911	OE1	GLU	A	114	-11.849	81.571	-56.581	1.00	31.54	T1
	ATOM	912	OE2	GLU	A	114	-10.008	82.780	-56.350	1.00	32.93	T1
	ATOM	913	C	GLU	A	114	-10.558	77.234	-54.899	1.00	29.39	T1
	ATOM	914	O	GLU	A	114	-9.846	76.734	-55.767	1.00	24.37	T1
	ATOM	915	N	GLY	A	115	-11.720	76.711	-54.540	1.00	33.97	T1
10	ATOM	916	CA	GLY	A	115	-12.185	75.509	-55.197	1.00	28.17	T1
	ATOM	917	C	GLY	A	115	-11.942	74.235	-54.426	1.00	24.33	T1
	ATOM	918	O	GLY	A	115	-12.562	73.212	-54.721	1.00	24.59	T1
	ATOM	919	N	ASP	A	116	-11.033	74.259	-53.458	1.00	30.73	T1
	ATOM	920	CA	ASP	A	116	-10.796	73.048	-52.688	1.00	31.24	T1
15	ATOM	921	CB	ASP	A	116	-9.613	73.211	-51.725	1.00	25.05	T1
	ATOM	922	CG	ASP	A	116	-8.270	73.264	-52.430	1.00	26.10	T1
	ATOM	923	OD1	ASP	A	116	-8.150	72.691	-53.534	1.00	24.18	T1
	ATOM	924	OD2	ASP	A	116	-7.327	73.861	-51.862	1.00	28.80	T1
	ATOM	925	C	ASP	A	116	-12.054	72.783	-51.878	1.00	27.72	T1
20	ATOM	926	O	ASP	A	116	-12.805	73.713	-51.558	1.00	33.19	T1
	ATOM	927	N	GLU	A	117	-12.288	71.519	-51.550	1.00	32.41	T1
	ATOM	928	CA	GLU	A	117	-13.450	71.169	-50.751	1.00	30.55	T1
	ATOM	929	CB	GLU	A	117	-14.446	70.359	-51.572	1.00	33.13	T1
	ATOM	930	CG	GLU	A	117	-14.867	71.025	-52.867	1.00	30.77	T1
25	ATOM	931	CD	GLU	A	117	-15.924	70.226	-53.616	1.00	27.00	T1
	ATOM	932	OE1	GLU	A	117	-15.849	68.969	-53.594	1.00	23.71	T1
	ATOM	933	OE2	GLU	A	117	-16.818	70.856	-54.234	1.00	26.98	T1
	ATOM	934	C	GLU	A	117	-12.988	70.351	-49.565	1.00	32.70	T1
	ATOM	935	O	GLU	A	117	-12.081	69.538	-49.690	1.00	22.77	T1
30	ATOM	936	N	LEU	A	118	-13.600	70.584	-48.412	1.00	33.21	T1
	ATOM	937	CA	LEU	A	118	-13.264	69.844	-47.206	1.00	34.28	T1
	ATOM	938	CB	LEU	A	118	-13.031	70.792	-46.032	1.00	36.99	T1
	ATOM	939	CG	LEU	A	118	-11.829	71.728	-46.124	1.00	21.88	T1
	ATOM	940	CD1	LEU	A	118	-11.722	72.530	-44.849	1.00	29.75	T1
35	ATOM	941	CD2	LEU	A	118	-10.568	70.919	-46.352	1.00	25.16	T1
	ATOM	942	C	LEU	A	118	-14.434	68.946	-46.878	1.00	31.75	T1
	ATOM	943	O	LEU	A	118	-15.586	69.311	-47.117	1.00	26.64	T1
	ATOM	944	N	GLN	A	119	-14.140	67.775	-46.329	1.00	31.38	T1
	ATOM	945	CA	GLN	A	119	-15.189	66.840	-45.950	1.00	30.58	T1
40	ATOM	946	CB	GLN	A	119	-15.568	65.978	-47.144	1.00	25.13	T1
	ATOM	947	CG	GLN	A	119	-14.453	65.090	-47.633	1.00	30.57	T1
	ATOM	948	CD	GLN	A	119	-14.842	64.305	-48.869	1.00	30.31	T1
	ATOM	949	OE1	GLN	A	119	-14.193	63.316	-49.220	1.00	29.37	T1
	ATOM	950	NE2	GLN	A	119	-15.899	64.750	-49.546	1.00	30.76	T1
45	ATOM	951	C	GLN	A	119	-14.764	65.962	-44.776	1.00	32.40	T1
	ATOM	952	O	GLN	A	119	-13.574	65.740	-44.561	1.00	24.95	T1
	ATOM	953	N	LEU	A	120	-15.749	65.478	-44.023	1.00	33.10	T1
	ATOM	954	CA	LEU	A	120	-15.525	64.633	-42.849	1.00	30.19	T1
	ATOM	955	CB	LEU	A	120	-16.346	65.180	-41.681	1.00	29.81	T1
50	ATOM	956	CG	LEU	A	120	-16.184	64.744	-40.222	1.00	26.82	T1
	ATOM	957	CD1	LEU	A	120	-16.100	63.239	-40.121	1.00	30.47	T1
	ATOM	958	CD2	LEU	A	120	-14.951	65.390	-39.642	1.00	30.80	T1
	ATOM	959	C	LEU	A	120	-16.007	63.234	-43.208	1.00	23.13	T1
	ATOM	960	O	LEU	A	120	-17.179	63.050	-43.536	1.00	25.85	T1
55	ATOM	961	N	ALA	A	121	-15.127	62.243	-43.143	1.00	28.50	T1
	ATOM	962	CA	ALA	A	121	-15.527	60.889	-43.508	1.00	26.21	T1
	ATOM	963	CB	ALA	A	121	-14.900	60.522	-44.840	1.00	26.86	T1
	ATOM	964	C	ALA	A	121	-15.220	59.797	-42.482	1.00	29.69	T1
	ATOM	965	O	ALA	A	121	-14.165	59.798	-41.841	1.00	25.25	T1
60	ATOM	966	N	ILE	A	122	-16.155	58.859	-42.337	1.00	28.00	T1
	ATOM	967	CA	ILE	A	122	-15.979	57.741	-41.420	1.00	35.42	T1
	ATOM	968	CB	ILE	A	122	-17.211	57.533	-40.539	1.00	28.40	T1
	ATOM	969	CG2	ILE	A	122	-16.959	56.388	-39.577	1.00	35.40	T1
	ATOM	970	CG1	ILE	A	122	-17.508	58.815	-39.759	1.00	37.48	T1
65	ATOM	971	CD1	ILE	A	122	-18.712	58.724	-38.853	1.00	27.05	T1
	ATOM	972	C	ILE	A	122	-15.743	56.484	-42.251	1.00	26.26	T1

	ATOM	973	O	ILE A 122	-16.602	56.071	-43.044	1.00	34.67	T1
	ATOM	974	N	PRO A 123	-14.563	55.860	-42.083	1.00	23.95	T1
	ATOM	975	CD	PRO A 123	-13.454	56.357	-41.258	1.00	26.48	T1
	ATOM	976	CA	PRO A 123	-14.160	54.641	-42.803	1.00	32.16	T1
5	ATOM	977	CB	PRO A 123	-12.663	54.532	-42.501	1.00	20.50	T1
	ATOM	978	CG	PRO A 123	-12.278	55.923	-42.074	1.00	26.71	T1
	ATOM	979	C	PRO A 123	-14.908	53.379	-42.363	1.00	30.95	T1
	ATOM	980	O	PRO A 123	-14.285	52.387	-41.981	1.00	28.06	T1
	ATOM	981	N	ARG A 124	-16.236	53.424	-42.399	1.00	27.49	T1
10	ATOM	982	CA	ARG A 124	-17.048	52.281	-42.023	1.00	29.86	T1
	ATOM	983	CB	ARG A 124	-17.400	52.314	-40.554	1.00	31.39	T1
	ATOM	984	CG	ARG A 124	-16.346	51.690	-39.684	1.00	27.02	T1
	ATOM	985	CD	ARG A 124	-17.022	50.829	-38.638	1.00	28.81	T1
	ATOM	986	NE	ARG A 124	-17.559	49.593	-39.209	1.00	21.80	T1
15	ATOM	987	CZ	ARG A 124	-18.348	48.755	-38.539	1.00	29.38	T1
	ATOM	988	NH1	ARG A 124	-18.691	49.036	-37.284	1.00	33.88	T1
	ATOM	989	NH2	ARG A 124	-18.779	47.625	-39.107	1.00	26.34	T1
	ATOM	990	C	ARG A 124	-18.312	52.296	-42.834	1.00	26.01	T1
	ATOM	991	O	ARG A 124	-18.734	53.358	-43.307	1.00	34.35	T1
20	ATOM	992	N	GLU A 125	-18.927	51.124	-42.986	1.00	26.83	T1
	ATOM	993	CA	GLU A 125	-20.133	51.025	-43.791	1.00	34.14	T1
	ATOM	994	CB	GLU A 125	-20.389	49.575	-44.178	1.00	34.44	T1
	ATOM	995	CG	GLU A 125	-19.395	49.082	-45.227	1.00	32.31	T1
	ATOM	996	CD	GLU A 125	-20.062	48.233	-46.311	1.00	39.35	T1
25	ATOM	997	OE1	GLU A 125	-20.726	47.219	-45.947	1.00	33.52	T1
	ATOM	998	OE2	GLU A 125	-19.915	48.579	-47.520	1.00	31.16	T1
	ATOM	999	C	GLU A 125	-21.374	51.650	-43.178	1.00	30.43	T1
	ATOM	1000	O	GLU A 125	-22.060	52.429	-43.843	1.00	33.01	T1
	ATOM	1001	N	ASN A 126	-21.685	51.316	-41.934	1.00	30.02	T1
30	ATOM	1002	CA	ASN A 126	-22.843	51.938	-41.316	1.00	27.89	T1
	ATOM	1003	CB	ASN A 126	-24.058	51.019	-41.341	1.00	27.38	T1
	ATOM	1004	CG	ASN A 126	-24.777	51.051	-42.686	1.00	33.64	T1
	ATOM	1005	OD1	ASN A 126	-24.340	50.420	-43.659	1.00	31.78	T1
	ATOM	1006	ND2	ASN A 126	-25.878	51.806	-42.752	1.00	30.28	T1
35	ATOM	1007	C	ASN A 126	-22.491	52.299	-39.908	1.00	29.09	T1
	ATOM	1008	O	ASN A 126	-23.065	51.775	-38.949	1.00	32.82	T1
	ATOM	1009	N	ALA A 127	-21.525	53.206	-39.802	1.00	28.21	T1
	ATOM	1010	CA	ALA A 127	-21.035	53.668	-38.517	1.00	32.43	T1
	ATOM	1011	CB	ALA A 127	-20.125	54.864	-38.726	1.00	27.52	T1
40	ATOM	1012	C	ALA A 127	-22.163	54.034	-37.567	1.00	31.71	T1
	ATOM	1013	O	ALA A 127	-23.097	54.741	-37.947	1.00	31.36	T1
	ATOM	1014	N	GLN A 128	-22.081	53.528	-36.341	1.00	24.27	T1
	ATOM	1015	CA	GLN A 128	-23.084	53.846	-35.323	1.00	25.61	T1
	ATOM	1016	CB	GLN A 128	-23.147	52.738	-34.266	1.00	28.10	T1
45	ATOM	1017	CG	GLN A 128	-23.673	51.431	-34.832	1.00	37.01	T1
	ATOM	1018	CD	GLN A 128	-24.923	51.652	-35.690	1.00	25.68	T1
	ATOM	1019	OE1	GLN A 128	-25.938	52.189	-35.220	1.00	27.48	T1
	ATOM	1020	NE2	GLN A 128	-24.849	51.246	-36.957	1.00	34.68	T1
	ATOM	1021	C	GLN A 128	-22.634	55.171	-34.707	1.00	28.43	T1
50	ATOM	1022	O	GLN A 128	-21.746	55.212	-33.842	1.00	32.54	T1
	ATOM	1023	N	ILE A 129	-23.273	56.243	-35.161	1.00	27.16	T1
	ATOM	1024	CA	ILE A 129	-22.923	57.599	-34.765	1.00	30.20	T1
	ATOM	1025	CB	ILE A 129	-22.422	58.352	-36.061	1.00	35.93	T1
	ATOM	1026	CG2	ILE A 129	-23.129	59.680	-36.262	1.00	28.56	T1
55	ATOM	1027	CG1	ILE A 129	-20.905	58.479	-36.019	1.00	28.41	T1
	ATOM	1028	CD1	ILE A 129	-20.204	57.137	-35.963	1.00	31.13	T1
	ATOM	1029	C	ILE A 129	-24.058	58.367	-34.095	1.00	30.32	T1
	ATOM	1030	O	ILE A 129	-25.228	58.012	-34.245	1.00	27.85	T1
	ATOM	1031	N	SER A 130	-23.712	59.405	-33.341	1.00	34.17	T1
60	ATOM	1032	CA	SER A 130	-24.729	60.240	-32.716	1.00	32.58	T1
	ATOM	1033	CB	SER A 130	-24.328	60.621	-31.291	1.00	31.18	T1
	ATOM	1034	OG	SER A 130	-25.184	61.639	-30.782	1.00	35.12	T1
	ATOM	1035	C	SER A 130	-24.843	61.505	-33.560	1.00	31.18	T1
	ATOM	1036	O	SER A 130	-23.845	62.170	-33.813	1.00	34.66	T1
65	ATOM	1037	N	LEU A 131	-26.045	61.845	-34.000	1.00	32.91	T1
	ATOM	1038	CA	LEU A 131	-26.213	63.040	-34.815	1.00	33.14	T1

	ATOM	1039	CB	LEU	A	131	-27.182	62.774	-35.967	1.00	35.13	T1
	ATOM	1040	CG	LEU	A	131	-26.596	62.022	-37.163	1.00	31.80	T1
	ATOM	1041	CD1	LEU	A	131	-26.065	60.669	-36.724	1.00	33.50	T1
	ATOM	1042	CD2	LEU	A	131	-27.666	61.856	-38.217	1.00	31.40	T1
5	ATOM	1043	C	LEU	A	131	-26.669	64.273	-34.055	1.00	28.73	T1
	ATOM	1044	O	LEU	A	131	-27.394	65.108	-34.599	1.00	25.72	T1
	ATOM	1045	N	ASP	A	132	-26.242	64.400	-32.805	1.00	33.94	T1
	ATOM	1046	CA	ASP	A	132	-26.615	65.559	-32.003	1.00	34.99	T1
	ATOM	1047	CB	ASP	A	132	-26.630	65.197	-30.515	1.00	27.15	T1
10	ATOM	1048	CG	ASP	A	132	-27.937	64.539	-30.086	1.00	35.81	T1
	ATOM	1049	OD1	ASP	A	132	-27.974	63.986	-28.960	1.00	32.83	T1
	ATOM	1050	OD2	ASP	A	132	-28.923	64.587	-30.869	1.00	29.22	T1
	ATOM	1051	C	ASP	A	132	-25.680	66.740	-32.236	1.00	22.15	T1
	ATOM	1052	O	ASP	A	132	-24.466	66.576	-32.401	1.00	30.71	T1
15	ATOM	1053	N	GLY	A	133	-26.263	67.933	-32.242	1.00	33.50	T1
	ATOM	1054	CA	GLY	A	133	-25.496	69.148	-32.447	1.00	26.34	T1
	ATOM	1055	C	GLY	A	133	-24.311	69.344	-31.518	1.00	25.91	T1
	ATOM	1056	O	GLY	A	133	-23.324	69.937	-31.932	1.00	30.09	T1
	ATOM	1057	N	ASP	A	134	-24.390	68.861	-30.282	1.00	31.40	T1
20	ATOM	1058	CA	ASP	A	134	-23.285	69.030	-29.348	1.00	39.11	T1
	ATOM	1059	CB	ASP	A	134	-23.623	68.591	-27.928	1.00	30.54	T1
	ATOM	1060	CG	ASP	A	134	-25.065	68.656	-27.618	1.00	31.61	T1
	ATOM	1061	OD1	ASP	A	134	-25.591	69.784	-27.641	1.00	31.72	T1
	ATOM	1062	OD2	ASP	A	134	-25.663	67.584	-27.343	1.00	36.95	T1
25	ATOM	1063	C	ASP	A	134	-22.120	68.160	-29.711	1.00	32.41	T1
	ATOM	1064	O	ASP	A	134	-21.003	68.627	-29.892	1.00	26.34	T1
	ATOM	1065	N	VAL	A	135	-22.393	66.867	-29.762	1.00	32.38	T1
	ATOM	1066	CA	VAL	A	135	-21.368	65.876	-30.010	1.00	30.77	T1
	ATOM	1067	CB	VAL	A	135	-21.911	64.513	-29.637	1.00	25.30	T1
30	ATOM	1068	CG1	VAL	A	135	-22.292	64.531	-28.163	1.00	27.77	T1
	ATOM	1069	CG2	VAL	A	135	-23.130	64.191	-30.489	1.00	27.81	T1
	ATOM	1070	C	VAL	A	135	-20.660	65.790	-31.359	1.00	31.01	T1
	ATOM	1071	O	VAL	A	135	-19.462	65.513	-31.383	1.00	33.63	T1
	ATOM	1072	N	THR	A	136	-21.349	66.006	-32.477	1.00	33.11	T1
35	ATOM	1073	CA	THR	A	136	-20.636	65.925	-33.742	1.00	27.96	T1
	ATOM	1074	CB	THR	A	136	-21.035	64.654	-34.531	1.00	35.02	T1
	ATOM	1075	OG1	THR	A	136	-22.304	64.844	-35.145	1.00	33.87	T1
	ATOM	1076	CG2	THR	A	136	-21.137	63.464	-33.607	1.00	39.89	T1
	ATOM	1077	C	THR	A	136	-20.794	67.178	-34.607	1.00	35.58	T1
40	ATOM	1078	O	THR	A	136	-21.901	67.576	-34.969	1.00	31.04	T1
	ATOM	1079	N	PHE	A	137	-19.662	67.803	-34.922	1.00	31.15	T1
	ATOM	1080	CA	PHE	A	137	-19.639	69.023	-35.727	1.00	31.35	T1
	ATOM	1081	CB	PHE	A	137	-19.714	70.230	-34.797	1.00	29.64	T1
	ATOM	1082	CG	PHE	A	137	-18.831	70.122	-33.588	1.00	28.13	T1
45	ATOM	1083	CD1	PHE	A	137	-17.476	70.412	-33.668	1.00	31.24	T1
	ATOM	1084	CD2	PHE	A	137	-19.354	69.732	-32.366	1.00	29.83	T1
	ATOM	1085	CE1	PHE	A	137	-16.663	70.318	-32.550	1.00	34.20	T1
	ATOM	1086	CE2	PHE	A	137	-18.546	69.634	-31.239	1.00	33.37	T1
	ATOM	1087	CZ	PHE	A	137	-17.204	69.927	-31.331	1.00	28.78	T1
50	ATOM	1088	C	PHE	A	137	-18.407	69.110	-36.636	1.00	26.37	T1
	ATOM	1089	O	PHE	A	137	-17.437	68.371	-36.448	1.00	31.37	T1
	ATOM	1090	N	PHE	A	138	-18.444	70.016	-37.611	1.00	25.28	T1
	ATOM	1091	CA	PHE	A	138	-17.344	70.158	-38.555	1.00	28.17	T1
	ATOM	1092	CB	PHE	A	138	-17.813	69.605	-39.901	1.00	30.10	T1
55	ATOM	1093	CG	PHE	A	138	-16.717	69.383	-40.898	1.00	31.35	T1
	ATOM	1094	CD1	PHE	A	138	-15.405	69.188	-40.488	1.00	33.10	T1
	ATOM	1095	CD2	PHE	A	138	-17.005	69.365	-42.262	1.00	31.59	T1
	ATOM	1096	CE1	PHE	A	138	-14.391	68.981	-41.421	1.00	32.37	T1
	ATOM	1097	CE2	PHE	A	138	-16.003	69.159	-43.198	1.00	25.63	T1
60	ATOM	1098	CZ	PHE	A	138	-14.692	68.968	-42.779	1.00	24.22	T1
	ATOM	1099	C	PHE	A	138	-16.834	71.612	-38.651	1.00	30.52	T1
	ATOM	1100	O	PHE	A	138	-17.607	72.554	-38.821	1.00	28.80	T1
	ATOM	1101	N	GLY	A	139	-15.508	71.750	-38.556	1.00	27.47	T1
	ATOM	1102	CA	GLY	A	139	-14.808	73.034	-38.535	1.00	31.91	T1
65	ATOM	1103	C	GLY	A	139	-14.594	73.997	-39.683	1.00	32.41	T1
	ATOM	1104	O	GLY	A	139	-15.530	74.311	-40.395	1.00	35.60	T1

	ATOM	1105	N	ALA	A	140	-13.364	74.505	-39.804	1.00	37.44	T1
	ATOM	1106	CA	ALA	A	140	-12.932	75.469	-40.841	1.00	28.74	T1
	ATOM	1107	CB	ALA	A	140	-13.632	75.193	-42.145	1.00	32.86	T1
	ATOM	1108	C	ALA	A	140	-13.054	76.966	-40.509	1.00	26.70	T1
5	ATOM	1109	O	ALA	A	140	-14.143	77.537	-40.522	1.00	34.03	T1
	ATOM	1110	N	LEU	A	141	-11.908	77.589	-40.236	1.00	33.69	T1
	ATOM	1111	CA	LEU	A	141	-11.811	79.015	-39.899	1.00	29.59	T1
	ATOM	1112	CB	LEU	A	141	-11.693	79.200	-38.373	1.00	26.29	T1
	ATOM	1113	CG	LEU	A	141	-11.409	80.595	-37.806	1.00	26.44	T1
10	ATOM	1114	CD1	LEU	A	141	-11.743	80.638	-36.345	1.00	31.81	T1
	ATOM	1115	CD2	LEU	A	141	-9.956	80.935	-38.000	1.00	30.16	T1
	ATOM	1116	C	LEU	A	141	-10.566	79.572	-40.585	1.00	27.97	T1
	ATOM	1117	O	LEU	A	141	-9.502	78.967	-40.533	1.00	30.17	T1
	ATOM	1118	N	LYS	A	142	-10.689	80.731	-41.213	1.00	29.79	T1
15	ATOM	1119	CA	LYS	A	142	-9.555	81.311	-41.916	1.00	23.67	T1
	ATOM	1120	CB	LYS	A	142	-10.042	82.118	-43.116	1.00	29.27	T1
	ATOM	1121	CG	LYS	A	142	-8.907	82.742	-43.890	1.00	24.33	T1
	ATOM	1122	CD	LYS	A	142	-9.328	83.283	-45.238	1.00	33.74	T1
	ATOM	1123	CE	LYS	A	142	-8.122	83.872	-45.962	1.00	27.23	T1
20	ATOM	1124	NZ	LYS	A	142	-8.489	84.450	-47.283	1.00	35.50	T1
	ATOM	1125	C	LYS	A	142	-8.627	82.181	-41.074	1.00	33.03	T1
	ATOM	1126	O	LYS	A	142	-9.071	83.100	-40.396	1.00	28.02	T1
	ATOM	1127	N	LEU	A	143	-7.329	81.894	-41.134	1.00	30.35	T1
	ATOM	1128	CA	LEU	A	143	-6.330	82.659	-40.386	1.00	28.94	T1
25	ATOM	1129	CB	LEU	A	143	-5.060	81.826	-40.189	1.00	25.75	T1
	ATOM	1130	CG	LEU	A	143	-5.169	80.473	-39.484	1.00	33.45	T1
	ATOM	1131	CD1	LEU	A	143	-3.814	79.784	-39.490	1.00	27.67	T1
	ATOM	1132	CD2	LEU	A	143	-5.657	80.671	-38.070	1.00	31.92	T1
	ATOM	1133	C	LEU	A	143	-5.968	83.937	-41.137	1.00	30.17	T1
30	ATOM	1134	O	LEU	A	143	-6.107	84.011	-42.358	1.00	24.33	T1
	ATOM	1135	N	LEU	A	144	-5.498	84.942	-40.406	1.00	30.48	T1
	ATOM	1136	CA	LEU	A	144	-5.106	86.196	-41.035	1.00	32.90	T1
	ATOM	1137	CB	LEU	A	144	-5.257	87.356	-40.058	1.00	36.81	T1
	ATOM	1138	CG	LEU	A	144	-6.688	87.671	-39.634	1.00	21.03	T1
35	ATOM	1139	CD1	LEU	A	144	-6.666	88.713	-38.531	1.00	27.39	T1
	ATOM	1140	CD2	LEU	A	144	-7.496	88.175	-40.830	1.00	28.98	T1
	ATOM	1141	C	LEU	A	144	-3.659	86.111	-41.492	1.00	26.60	T1
	ATOM	1142	O	LEU	A	144	-2.929	85.213	-41.013	1.00	32.79	T1
	ATOM	1143	OXT	LEU	A	144	-3.272	86.959	-42.320	1.00	30.28	T1
40	ATOM	1144	CB	VAL	A	1	-0.327	92.342	-41.378	1.00	32.68	T2
	ATOM	1145	CG1	VAL	A	1	-1.097	92.176	-42.700	1.00	23.07	T2
	ATOM	1146	CG2	VAL	A	1	0.133	93.789	-41.216	1.00	29.26	T2
	ATOM	1147	C	VAL	A	1	-1.572	90.425	-40.290	1.00	31.27	T2
	ATOM	1148	O	VAL	A	1	-2.750	90.076	-40.411	1.00	31.11	T2
45	ATOM	1149	N	VAL	A	1	-0.565	92.226	-38.861	1.00	29.81	T2
	ATOM	1150	CA	VAL	A	1	-1.226	91.921	-40.165	1.00	34.24	T2
	ATOM	1151	N	THR	A	2	-0.567	89.544	-40.260	1.00	33.96	T2
	ATOM	1152	CA	THR	A	2	-0.831	88.110	-40.383	1.00	30.04	T2
	ATOM	1153	CB	THR	A	2	-0.023	87.482	-41.523	1.00	28.70	T2
50	ATOM	1154	OG1	THR	A	2	1.356	87.427	-41.147	1.00	38.22	T2
	ATOM	1155	CG2	THR	A	2	-0.177	88.295	-42.800	1.00	31.82	T2
	ATOM	1156	C	THR	A	2	-0.518	87.328	-39.113	1.00	34.46	T2
	ATOM	1157	O	THR	A	2	0.051	87.866	-38.171	1.00	28.76	T2
	ATOM	1158	N	GLN	A	3	-0.884	86.049	-39.103	1.00	31.48	T2
55	ATOM	1159	CA	GLN	A	3	-0.645	85.189	-37.947	1.00	31.47	T2
	ATOM	1160	CB	GLN	A	3	-1.948	84.521	-37.505	1.00	31.69	T2
	ATOM	1161	CG	GLN	A	3	-3.135	85.454	-37.416	1.00	31.84	T2
	ATOM	1162	CD	GLN	A	3	-4.348	84.771	-36.836	1.00	29.04	T2
	ATOM	1163	OE1	GLN	A	3	-4.312	84.301	-35.704	1.00	30.60	T2
60	ATOM	1164	NE2	GLN	A	3	-5.427	84.705	-37.608	1.00	33.66	T2
	ATOM	1165	C	GLN	A	3	0.371	84.096	-38.256	1.00	31.36	T2
	ATOM	1166	O	GLN	A	3	0.047	83.111	-38.931	1.00	35.58	T2
	ATOM	1167	N	ASP	A	4	1.594	84.253	-37.757	1.00	26.40	T2
	ATOM	1168	CA	ASP	A	4	2.611	83.242	-37.999	1.00	32.70	T2
65	ATOM	1169	CB	ASP	A	4	3.947	83.666	-37.398	1.00	32.73	T2
	ATOM	1170	CG	ASP	A	4	4.503	84.922	-38.040	1.00	26.60	T2

	ATOM	1171	OD1	ASP	A	4	4.059	85.257	-39.158	1.00	25.93	T2
	ATOM	1172	OD2	ASP	A	4	5.396	85.567	-37.439	1.00	25.56	T2
	ATOM	1173	C	ASP	A	4	2.184	81.916	-37.394	1.00	25.39	T2
	ATOM	1174	O	ASP	A	4	1.502	81.881	-36.372	1.00	31.34	T2
5	ATOM	1175	N	CYS	A	5	2.576	80.828	-38.041	1.00	35.20	T2
	ATOM	1176	CA	CYS	A	5	2.258	79.484	-37.573	1.00	22.44	T2
	ATOM	1177	CB	CYS	A	5	0.830	79.078	-37.990	1.00	32.34	T2
	ATOM	1178	SG	CYS	A	5	0.199	79.805	-39.531	1.00	36.81	T2
	ATOM	1179	C	CYS	A	5	3.271	78.481	-38.120	1.00	36.01	T2
10	ATOM	1180	O	CYS	A	5	3.803	78.650	-39.212	1.00	32.08	T2
	ATOM	1181	N	LEU	A	6	3.567	77.448	-37.344	1.00	37.97	T2
	ATOM	1182	CA	LEU	A	6	4.500	76.421	-37.783	1.00	30.55	T2
	ATOM	1183	CB	LEU	A	6	5.870	76.630	-37.142	1.00	36.56	T2
	ATOM	1184	CG	LEU	A	6	6.950	75.590	-37.448	1.00	31.86	T2
15	ATOM	1185	CD1	LEU	A	6	8.312	76.229	-37.384	1.00	32.24	T2
	ATOM	1186	CD2	LEU	A	6	6.862	74.452	-36.453	1.00	24.58	T2
	ATOM	1187	C	LEU	A	6	3.914	75.093	-37.357	1.00	31.00	T2
	ATOM	1188	O	LEU	A	6	3.450	74.955	-36.233	1.00	34.34	T2
	ATOM	1189	N	GLN	A	7	3.926	74.120	-38.258	1.00	34.18	T2
20	ATOM	1190	CA	GLN	A	7	3.363	72.814	-37.950	1.00	32.87	T2
	ATOM	1191	CB	GLN	A	7	2.058	72.623	-38.712	1.00	27.12	T2
	ATOM	1192	CG	GLN	A	7	1.250	71.425	-38.273	1.00	38.89	T2
	ATOM	1193	CD	GLN	A	7	-0.118	71.431	-38.894	1.00	26.22	T2
	ATOM	1194	OE1	GLN	A	7	-0.258	71.240	-40.093	1.00	27.86	T2
25	ATOM	1195	NE2	GLN	A	7	-1.138	71.676	-38.086	1.00	29.14	T2
	ATOM	1196	C	GLN	A	7	4.313	71.681	-38.285	1.00	29.78	T2
	ATOM	1197	O	GLN	A	7	5.005	71.716	-39.303	1.00	32.37	T2
	ATOM	1198	N	LEU	A	8	4.331	70.674	-37.418	1.00	29.67	T2
	ATOM	1199	CA	LEU	A	8	5.193	69.520	-37.600	1.00	25.35	T2
30	ATOM	1200	CB	LEU	A	8	6.127	69.367	-36.389	1.00	29.67	T2
	ATOM	1201	CG	LEU	A	8	7.462	70.119	-36.318	1.00	31.99	T2
	ATOM	1202	CD1	LEU	A	8	7.524	71.221	-37.342	1.00	27.29	T2
	ATOM	1203	CD2	LEU	A	8	7.633	70.662	-34.927	1.00	27.62	T2
	ATOM	1204	C	LEU	A	8	4.393	68.238	-37.805	1.00	31.41	T2
35	ATOM	1205	O	LEU	A	8	3.261	68.115	-37.347	1.00	35.35	T2
	ATOM	1206	N	ILE	A	9	5.009	67.284	-38.492	1.00	35.76	T2
	ATOM	1207	CA	ILE	A	9	4.395	65.993	-38.799	1.00	28.37	T2
	ATOM	1208	CB	ILE	A	9	4.140	65.864	-40.306	1.00	33.02	T2
	ATOM	1209	CG2	ILE	A	9	3.954	64.422	-40.687	1.00	31.69	T2
40	ATOM	1210	CG1	ILE	A	9	2.913	66.660	-40.710	1.00	22.57	T2
	ATOM	1211	CD1	ILE	A	9	2.675	66.582	-42.206	1.00	30.76	T2
	ATOM	1212	C	ILE	A	9	5.352	64.871	-38.412	1.00	30.39	T2
	ATOM	1213	O	ILE	A	9	6.568	65.009	-38.562	1.00	27.25	T2
	ATOM	1214	N	ALA	A	10	4.813	63.752	-37.940	1.00	35.56	T2
45	ATOM	1215	CA	ALA	A	10	5.669	62.635	-37.562	1.00	32.76	T2
	ATOM	1216	CB	ALA	A	10	4.845	61.512	-36.976	1.00	25.17	T2
	ATOM	1217	C	ALA	A	10	6.446	62.124	-38.766	1.00	26.74	T2
	ATOM	1218	O	ALA	A	10	5.876	61.894	-39.835	1.00	34.32	T2
	ATOM	1219	N	ASP	A	11	7.754	61.952	-38.590	1.00	33.58	T2
50	ATOM	1220	CA	ASP	A	11	8.604	61.447	-39.660	1.00	30.11	T2
	ATOM	1221	CB	ASP	A	11	10.000	62.059	-39.584	1.00	25.41	T2
	ATOM	1222	CG	ASP	A	11	10.943	61.452	-40.595	1.00	33.41	T2
	ATOM	1223	OD1	ASP	A	11	10.486	61.183	-41.728	1.00	33.77	T2
	ATOM	1224	OD2	ASP	A	11	12.132	61.250	-40.259	1.00	29.32	T2
55	ATOM	1225	C	ASP	A	11	8.698	59.935	-39.558	1.00	23.73	T2
	ATOM	1226	O	ASP	A	11	9.567	59.392	-38.877	1.00	29.05	T2
	ATOM	1227	N	SER	A	12	7.777	59.277	-40.251	1.00	34.28	T2
	ATOM	1228	CA	SER	A	12	7.670	57.824	-40.281	1.00	29.92	T2
	ATOM	1229	CB	SER	A	12	6.409	57.404	-41.064	1.00	35.65	T2
60	ATOM	1230	OG	SER	A	12	6.414	57.888	-42.405	1.00	25.68	T2
	ATOM	1231	C	SER	A	12	8.880	57.141	-40.897	1.00	25.93	T2
	ATOM	1232	O	SER	A	12	8.775	56.013	-41.372	1.00	25.86	T2
	ATOM	1233	N	GLU	A	13	10.027	57.813	-40.906	1.00	27.61	T2
	ATOM	1234	CA	GLU	A	13	11.212	57.198	-41.482	1.00	25.18	T2
65	ATOM	1235	CB	GLU	A	13	11.419	57.685	-42.903	1.00	32.04	T2
	ATOM	1236	CG	GLU	A	13	10.697	56.811	-43.890	1.00	29.74	T2

	ATOM	1237	CD	GLU	A	13	10.986	57.224	-45.313	1.00	28.74	T2
	ATOM	1238	OE1	GLU	A	13	12.174	57.535	-45.602	1.00	39.63	T2
	ATOM	1239	OE2	GLU	A	13	10.037	57.235	-46.147	1.00	33.79	T2
	ATOM	1240	C	GLU	A	13	12.494	57.337	-40.694	1.00	27.82	T2
5	ATOM	1241	O	GLU	A	13	13.590	57.279	-41.256	1.00	30.07	T2
	ATOM	1242	N	THR	A	14	12.347	57.533	-39.392	1.00	31.01	T2
	ATOM	1243	CA	THR	A	14	13.488	57.618	-38.502	1.00	29.38	T2
	ATOM	1244	CB	THR	A	14	14.002	59.062	-38.303	1.00	34.36	T2
	ATOM	1245	OG1	THR	A	14	12.938	59.882	-37.818	1.00	32.82	T2
10	ATOM	1246	CG2	THR	A	14	14.536	59.632	-39.611	1.00	27.17	T2
	ATOM	1247	C	THR	A	14	12.988	57.061	-37.184	1.00	30.13	T2
	ATOM	1248	O	THR	A	14	11.794	57.138	-36.869	1.00	29.63	T2
	ATOM	1249	N	PRO	A	15	13.894	56.466	-36.403	1.00	35.72	T2
	ATOM	1250	CD	PRO	A	15	15.333	56.341	-36.686	1.00	32.69	T2
15	ATOM	1251	CA	PRO	A	15	13.560	55.874	-35.102	1.00	25.30	T2
	ATOM	1252	CB	PRO	A	15	14.891	55.290	-34.629	1.00	39.82	T2
	ATOM	1253	CG	PRO	A	15	15.680	55.102	-35.912	1.00	28.53	T2
	ATOM	1254	C	PRO	A	15	13.045	56.929	-34.137	1.00	35.97	T2
	ATOM	1255	O	PRO	A	15	13.492	58.078	-34.162	1.00	31.10	T2
20	ATOM	1256	N	THR	A	16	12.112	56.538	-33.284	1.00	30.65	T2
	ATOM	1257	CA	THR	A	16	11.573	57.469	-32.310	1.00	33.55	T2
	ATOM	1258	CB	THR	A	16	10.352	56.869	-31.642	1.00	33.91	T2
	ATOM	1259	OG1	THR	A	16	10.756	55.778	-30.805	1.00	30.71	T2
	ATOM	1260	CG2	THR	A	16	9.403	56.343	-32.699	1.00	27.45	T2
25	ATOM	1261	C	THR	A	16	12.648	57.729	-31.259	1.00	28.57	T2
	ATOM	1262	O	THR	A	16	13.050	56.814	-30.544	1.00	32.82	T2
	ATOM	1263	N	ILE	A	17	13.108	58.971	-31.162	1.00	29.36	T2
	ATOM	1264	CA	ILE	A	17	14.147	59.331	-30.203	1.00	27.75	T2
	ATOM	1265	CB	ILE	A	17	14.382	60.846	-30.212	1.00	33.22	T2
30	ATOM	1266	CG2	ILE	A	17	15.455	61.217	-29.204	1.00	31.65	T2
	ATOM	1267	CG1	ILE	A	17	14.807	61.290	-31.615	1.00	35.60	T2
	ATOM	1268	CD1	ILE	A	17	15.010	62.784	-31.754	1.00	28.08	T2
	ATOM	1269	C	ILE	A	17	13.893	58.894	-28.758	1.00	34.77	T2
	ATOM	1270	O	ILE	A	17	12.813	59.123	-28.206	1.00	30.79	T2
35	ATOM	1271	N	GLN	A	18	14.904	58.264	-28.156	1.00	31.51	T2
	ATOM	1272	CA	GLN	A	18	14.825	57.803	-26.772	1.00	28.15	T2
	ATOM	1273	CB	GLN	A	18	15.151	56.323	-26.683	1.00	34.37	T2
	ATOM	1274	CG	GLN	A	18	14.055	55.539	-26.023	1.00	27.28	T2
	ATOM	1275	CD	GLN	A	18	12.860	55.391	-26.933	1.00	34.14	T2
40	ATOM	1276	OE1	GLN	A	18	12.867	54.578	-27.856	1.00	36.63	T2
	ATOM	1277	NE2	GLN	A	18	11.826	56.192	-26.691	1.00	26.04	T2
	ATOM	1278	C	GLN	A	18	15.794	58.579	-25.876	1.00	28.48	T2
	ATOM	1279	O	GLN	A	18	16.964	58.774	-26.231	1.00	32.65	T2
	ATOM	1280	N	LYS	A	19	15.316	59.011	-24.711	1.00	31.70	T2
45	ATOM	1281	CA	LYS	A	19	16.162	59.779	-23.802	1.00	29.09	T2
	ATOM	1282	CB	LYS	A	19	16.533	61.114	-24.452	1.00	23.95	T2
	ATOM	1283	CG	LYS	A	19	17.182	62.098	-23.504	1.00	39.47	T2
	ATOM	1284	CD	LYS	A	19	17.709	63.309	-24.264	1.00	31.42	T2
	ATOM	1285	CE	LYS	A	19	18.372	64.323	-23.317	1.00	33.32	T2
50	ATOM	1286	NZ	LYS	A	19	18.955	65.513	-24.029	1.00	27.90	T2
	ATOM	1287	C	LYS	A	19	15.506	60.032	-22.447	1.00	29.63	T2
	ATOM	1288	O	LYS	A	19	14.362	60.486	-22.374	1.00	37.71	T2
	ATOM	1289	N	GLY	A	20	16.249	59.747	-21.377	1.00	35.26	T2
	ATOM	1290	CA	GLY	A	20	15.733	59.934	-20.032	1.00	30.01	T2
55	ATOM	1291	C	GLY	A	20	14.433	59.178	-19.863	1.00	37.83	T2
	ATOM	1292	O	GLY	A	20	13.495	59.692	-19.247	1.00	32.75	T2
	ATOM	1293	N	SER	A	21	14.383	57.961	-20.414	1.00	26.38	T2
	ATOM	1294	CA	SER	A	21	13.188	57.100	-20.364	1.00	29.43	T2
	ATOM	1295	CB	SER	A	21	12.991	56.532	-18.946	1.00	24.20	T2
60	ATOM	1296	OG	SER	A	21	12.786	57.562	-17.986	1.00	38.52	T2
	ATOM	1297	C	SER	A	21	11.910	57.830	-20.834	1.00	32.97	T2
	ATOM	1298	O	SER	A	21	10.820	57.666	-20.261	1.00	37.24	T2
	ATOM	1299	N	TYR	A	22	12.079	58.639	-21.879	1.00	36.60	T2
	ATOM	1300	CA	TYR	A	22	11.004	59.407	-22.496	1.00	36.35	T2
65	ATOM	1301	CB	TYR	A	22	11.220	60.900	-22.274	1.00	31.93	T2
	ATOM	1302	CG	TYR	A	22	10.484	61.469	-21.093	1.00	31.49	T2

	ATOM	1303	CD1	TYR	A	22	10.116	60.663	-20.013	1.00	29.09	T2
	ATOM	1304	CE1	TYR	A	22	9.459	61.196	-18.902	1.00	28.21	T2
	ATOM	1305	CD2	TYR	A	22	10.177	62.825	-21.035	1.00	36.96	T2
	ATOM	1306	CE2	TYR	A	22	9.522	63.371	-19.927	1.00	28.20	T2
5	ATOM	1307	CZ	TYR	A	22	9.166	62.549	-18.863	1.00	33.58	T2
	ATOM	1308	OH	TYR	A	22	8.532	63.081	-17.763	1.00	32.15	T2
	ATOM	1309	C	TYR	A	22	11.081	59.122	-23.982	1.00	32.61	T2
	ATOM	1310	O	TYR	A	22	12.177	58.908	-24.520	1.00	30.37	T2
	ATOM	1311	N	THR	A	23	9.937	59.103	-24.654	1.00	26.17	T2
10	ATOM	1312	CA	THR	A	23	9.954	58.858	-26.089	1.00	32.16	T2
	ATOM	1313	CB	THR	A	23	8.927	57.793	-26.511	1.00	31.88	T2
	ATOM	1314	OG1	THR	A	23	8.914	56.716	-25.562	1.00	29.09	T2
	ATOM	1315	CG2	THR	A	23	9.308	57.236	-27.861	1.00	31.87	T2
	ATOM	1316	C	THR	A	23	9.630	60.165	-26.794	1.00	25.27	T2
15	ATOM	1317	O	THR	A	23	8.673	60.856	-26.436	1.00	37.06	T2
	ATOM	1318	N	PHE	A	24	10.446	60.516	-27.779	1.00	26.24	T2
	ATOM	1319	CA	PHE	A	24	10.232	61.742	-28.531	1.00	28.73	T2
	ATOM	1320	CB	PHE	A	24	11.440	62.666	-28.386	1.00	25.04	T2
	ATOM	1321	CG	PHE	A	24	11.641	63.183	-26.993	1.00	26.23	T2
20	ATOM	1322	CD1	PHE	A	24	12.249	62.395	-26.027	1.00	25.51	T2
	ATOM	1323	CD2	PHE	A	24	11.197	64.454	-26.641	1.00	27.43	T2
	ATOM	1324	CE1	PHE	A	24	12.411	62.867	-24.729	1.00	26.99	T2
	ATOM	1325	CE2	PHE	A	24	11.353	64.929	-25.349	1.00	26.86	T2
	ATOM	1326	CZ	PHE	A	24	11.960	64.135	-24.392	1.00	32.23	T2
25	ATOM	1327	C	PHE	A	24	9.969	61.467	-30.010	1.00	33.48	T2
	ATOM	1328	O	PHE	A	24	10.768	60.811	-30.693	1.00	26.95	T2
	ATOM	1329	N	VAL	A	25	8.841	61.975	-30.497	1.00	35.37	T2
	ATOM	1330	CA	VAL	A	25	8.459	61.798	-31.892	1.00	31.35	T2
	ATOM	1331	CB	VAL	A	25	7.052	62.357	-32.158	1.00	24.70	T2
30	ATOM	1332	CG1	VAL	A	25	6.708	62.225	-33.630	1.00	28.43	T2
	ATOM	1333	CG2	VAL	A	25	6.038	61.635	-31.299	1.00	29.71	T2
	ATOM	1334	C	VAL	A	25	9.431	62.509	-32.826	1.00	33.14	T2
	ATOM	1335	O	VAL	A	25	9.801	63.656	-32.598	1.00	35.89	T2
	ATOM	1336	N	PRO	A	26	9.874	61.822	-33.884	1.00	36.08	T2
35	ATOM	1337	CD	PRO	A	26	9.669	60.386	-34.138	1.00	29.57	T2
	ATOM	1338	CA	PRO	A	26	10.805	62.398	-34.861	1.00	32.79	T2
	ATOM	1339	CB	PRO	A	26	11.281	61.180	-35.652	1.00	27.79	T2
	ATOM	1340	CG	PRO	A	26	10.969	60.002	-34.749	1.00	30.08	T2
	ATOM	1341	C	PRO	A	26	9.988	63.336	-35.747	1.00	31.49	T2
40	ATOM	1342	O	PRO	A	26	9.145	62.872	-36.520	1.00	28.35	T2
	ATOM	1343	N	TRP	A	27	10.219	64.639	-35.654	1.00	24.38	T2
	ATOM	1344	CA	TRP	A	27	9.435	65.565	-36.461	1.00	31.85	T2
	ATOM	1345	CB	TRP	A	27	9.263	66.896	-35.728	1.00	32.48	T2
	ATOM	1346	CG	TRP	A	27	8.499	66.769	-34.463	1.00	27.28	T2
45	ATOM	1347	CD2	TRP	A	27	7.198	66.193	-34.297	1.00	34.22	T2
	ATOM	1348	CE2	TRP	A	27	6.896	66.237	-32.923	1.00	26.66	T2
	ATOM	1349	CE3	TRP	A	27	6.261	65.640	-35.175	1.00	32.67	T2
	ATOM	1350	CD1	TRP	A	27	8.918	67.135	-33.223	1.00	29.03	T2
	ATOM	1351	NE1	TRP	A	27	7.963	66.818	-32.288	1.00	29.17	T2
50	ATOM	1352	CZ2	TRP	A	27	5.695	65.750	-32.403	1.00	30.49	T2
	ATOM	1353	CZ3	TRP	A	27	5.065	65.154	-34.658	1.00	27.85	T2
	ATOM	1354	CH2	TRP	A	27	4.796	65.212	-33.285	1.00	26.97	T2
	ATOM	1355	C	TRP	A	27	9.947	65.831	-37.867	1.00	33.17	T2
	ATOM	1356	O	TRP	A	27	11.095	65.543	-38.207	1.00	29.49	T2
55	ATOM	1357	N	LEU	A	28	9.055	66.389	-38.676	1.00	28.88	T2
	ATOM	1358	CA	LEU	A	28	9.335	66.737	-40.055	1.00	30.14	T2
	ATOM	1359	CB	LEU	A	28	8.968	65.569	-40.959	1.00	35.78	T2
	ATOM	1360	CG	LEU	A	28	9.622	65.604	-42.335	1.00	34.47	T2
	ATOM	1361	CD1	LEU	A	28	11.143	65.503	-42.167	1.00	24.34	T2
60	ATOM	1362	CD2	LEU	A	28	9.080	64.458	-43.181	1.00	26.64	T2
	ATOM	1363	C	LEU	A	28	8.449	67.940	-40.361	1.00	24.64	T2
	ATOM	1364	O	LEU	A	28	7.244	67.913	-40.107	1.00	33.53	T2
	ATOM	1365	N	LEU	A	29	9.039	69.000	-40.896	1.00	34.74	T2
	ATOM	1366	CA	LEU	A	29	8.270	70.198	-41.187	1.00	31.10	T2
65	ATOM	1367	CB	LEU	A	29	9.145	71.252	-41.854	1.00	32.80	T2
	ATOM	1368	CG	LEU	A	29	8.369	72.515	-42.200	1.00	29.96	T2

	ATOM	1369	CD1	LEU	A	29	8.137	73.324	-40.939	1.00	24.20	T2
	ATOM	1370	CD2	LEU	A	29	9.127	73.317	-43.212	1.00	26.09	T2
	ATOM	1371	C	LEU	A	29	7.069	69.939	-42.076	1.00	29.97	T2
	ATOM	1372	O	LEU	A	29	7.192	69.364	-43.151	1.00	30.87	T2
5	ATOM	1373	N	SER	A	30	5.900	70.358	-41.607	1.00	30.16	T2
	ATOM	1374	CA	SER	A	30	4.680	70.223	-42.385	1.00	30.62	T2
	ATOM	1375	CB	SER	A	30	3.457	70.151	-41.483	1.00	30.16	T2
	ATOM	1376	OG	SER	A	30	2.279	70.125	-42.266	1.00	40.33	T2
	ATOM	1377	C	SER	A	30	4.623	71.497	-43.201	1.00	30.36	T2
10	ATOM	1378	O	SER	A	30	4.527	71.460	-44.415	1.00	30.71	T2
	ATOM	1379	N	PHE	A	31	4.692	72.629	-42.514	1.00	29.75	T2
	ATOM	1380	CA	PHE	A	31	4.686	73.925	-43.170	1.00	36.08	T2
	ATOM	1381	CB	PHE	A	31	3.299	74.222	-43.751	1.00	33.26	T2
	ATOM	1382	CG	PHE	A	31	2.352	74.866	-42.785	1.00	27.18	T2
15	ATOM	1383	CD1	PHE	A	31	2.372	76.240	-42.582	1.00	31.76	T2
	ATOM	1384	CD2	PHE	A	31	1.451	74.099	-42.067	1.00	31.67	T2
	ATOM	1385	CE1	PHE	A	31	1.507	76.841	-41.675	1.00	36.68	T2
	ATOM	1386	CE2	PHE	A	31	0.582	74.692	-41.158	1.00	32.44	T2
	ATOM	1387	CZ	PHE	A	31	0.611	76.069	-40.961	1.00	32.05	T2
20	ATOM	1388	C	PHE	A	31	5.089	74.990	-42.160	1.00	29.83	T2
	ATOM	1389	O	PHE	A	31	4.904	74.818	-40.949	1.00	27.69	T2
	ATOM	1390	N	LYS	A	32	5.658	76.081	-42.655	1.00	32.63	T2
	ATOM	1391	CA	LYS	A	32	6.070	77.177	-41.790	1.00	24.62	T2
	ATOM	1392	CB	LYS	A	32	7.587	77.174	-41.610	1.00	31.23	T2
25	ATOM	1393	CG	LYS	A	32	8.097	78.436	-40.973	1.00	32.96	T2
	ATOM	1394	CD	LYS	A	32	9.587	78.526	-41.005	1.00	23.91	T2
	ATOM	1395	CE	LYS	A	32	10.005	79.927	-40.616	1.00	32.56	T2
	ATOM	1396	NZ	LYS	A	32	11.476	80.044	-40.469	1.00	31.17	T2
	ATOM	1397	C	LYS	A	32	5.619	78.489	-42.418	1.00	27.88	T2
30	ATOM	1398	O	LYS	A	32	5.911	78.769	-43.575	1.00	35.73	T2
	ATOM	1399	N	ARG	A	33	4.907	79.294	-41.648	1.00	31.58	T2
	ATOM	1400	CA	ARG	A	33	4.394	80.562	-42.148	1.00	27.08	T2
	ATOM	1401	CB	ARG	A	33	2.870	80.460	-42.288	1.00	34.14	T2
	ATOM	1402	CG	ARG	A	33	2.135	81.730	-42.638	1.00	34.66	T2
35	ATOM	1403	CD	ARG	A	33	0.772	81.367	-43.210	1.00	26.21	T2
	ATOM	1404	NE	ARG	A	33	-0.064	82.531	-43.506	1.00	38.13	T2
	ATOM	1405	CZ	ARG	A	33	-0.794	83.173	-42.597	1.00	24.69	T2
	ATOM	1406	NH1	ARG	A	33	-0.798	82.758	-41.332	1.00	25.60	T2
	ATOM	1407	NH2	ARG	A	33	-1.508	84.236	-42.946	1.00	28.58	T2
40	ATOM	1408	C	ARG	A	33	4.770	81.699	-41.204	1.00	31.79	T2
	ATOM	1409	O	ARG	A	33	4.457	81.671	-40.008	1.00	29.79	T2
	ATOM	1410	N	GLY	A	34	5.455	82.699	-41.743	1.00	31.43	T2
	ATOM	1411	CA	GLY	A	34	5.855	83.820	-40.918	1.00	29.76	T2
	ATOM	1412	C	GLY	A	34	7.249	83.677	-40.346	1.00	27.75	T2
45	ATOM	1413	O	GLY	A	34	8.036	82.831	-40.773	1.00	34.55	T2
	ATOM	1414	N	SER	A	35	7.546	84.498	-39.350	1.00	34.22	T2
	ATOM	1415	CA	SER	A	35	8.859	84.494	-38.729	1.00	27.97	T2
	ATOM	1416	CB	SER	A	35	9.509	85.853	-38.948	1.00	32.07	T2
	ATOM	1417	OG	SER	A	35	8.658	86.874	-38.463	1.00	30.81	T2
50	ATOM	1418	C	SER	A	35	8.894	84.174	-37.236	1.00	32.97	T2
	ATOM	1419	O	SER	A	35	9.937	83.783	-36.717	1.00	37.23	T2
	ATOM	1420	N	ALA	A	36	7.771	84.331	-36.546	1.00	29.59	T2
	ATOM	1421	CA	ALA	A	36	7.733	84.097	-35.108	1.00	33.33	T2
	ATOM	1422	CB	ALA	A	36	6.379	84.502	-34.562	1.00	30.06	T2
55	ATOM	1423	C	ALA	A	36	8.076	82.697	-34.626	1.00	21.84	T2
	ATOM	1424	O	ALA	A	36	8.365	82.518	-33.446	1.00	35.20	T2
	ATOM	1425	N	LEU	A	37	8.050	81.707	-35.516	1.00	27.84	T2
	ATOM	1426	CA	LEU	A	37	8.351	80.330	-35.114	1.00	27.90	T2
	ATOM	1427	CB	LEU	A	37	7.045	79.544	-34.971	1.00	28.29	T2
60	ATOM	1428	CG	LEU	A	37	6.079	80.112	-33.927	1.00	25.03	T2
	ATOM	1429	CD1	LEU	A	37	4.687	79.580	-34.158	1.00	29.55	T2
	ATOM	1430	CD2	LEU	A	37	6.570	79.756	-32.547	1.00	29.90	T2
	ATOM	1431	C	LEU	A	37	9.294	79.609	-36.074	1.00	29.18	T2
	ATOM	1432	O	LEU	A	37	9.274	79.860	-37.274	1.00	35.64	T2
65	ATOM	1433	N	GLU	A	38	10.114	78.710	-35.533	1.00	27.22	T2
	ATOM	1434	CA	GLU	A	38	11.087	77.956	-36.325	1.00	29.93	T2

	ATOM	1435	CB	GLU	A	38	12.443	78.660	-36.304	1.00	27.09	T2
	ATOM	1436	CG	GLU	A	38	12.606	79.800	-37.284	1.00	29.58	T2
	ATOM	1437	CD	GLU	A	38	13.873	80.602	-37.018	1.00	34.34	T2
	ATOM	1438	OE1	GLU	A	38	14.886	79.984	-36.615	1.00	33.57	T2
5	ATOM	1439	OE2	GLU	A	38	13.854	81.842	-37.218	1.00	40.04	T2
	ATOM	1440	C	GLU	A	38	11.302	76.548	-35.789	1.00	27.95	T2
	ATOM	1441	O	GLU	A	38	10.934	76.250	-34.657	1.00	34.02	T2
	ATOM	1442	N	GLU	A	39	11.904	75.685	-36.606	1.00	34.47	T2
	ATOM	1443	CA	GLU	A	39	12.212	74.326	-36.168	1.00	29.60	T2
10	ATOM	1444	CB	GLU	A	39	12.270	73.340	-37.317	1.00	34.45	T2
	ATOM	1445	CG	GLU	A	39	11.203	73.457	-38.334	1.00	35.74	T2
	ATOM	1446	CD	GLU	A	39	11.707	72.990	-39.685	1.00	27.05	T2
	ATOM	1447	OE1	GLU	A	39	12.117	73.864	-40.492	1.00	26.71	T2
	ATOM	1448	OE2	GLU	A	39	11.720	71.752	-39.931	1.00	27.45	T2
15	ATOM	1449	C	GLU	A	39	13.620	74.401	-35.633	1.00	31.60	T2
	ATOM	1450	O	GLU	A	39	14.447	75.152	-36.150	1.00	31.13	T2
	ATOM	1451	N	LYS	A	40	13.912	73.616	-34.612	1.00	31.80	T2
	ATOM	1452	CA	LYS	A	40	15.261	73.606	-34.076	1.00	27.18	T2
	ATOM	1453	CB	LYS	A	40	15.507	74.823	-33.193	1.00	28.70	T2
20	ATOM	1454	CG	LYS	A	40	16.923	74.877	-32.653	1.00	31.71	T2
	ATOM	1455	CD	LYS	A	40	17.029	75.847	-31.488	1.00	24.40	T2
	ATOM	1456	CE	LYS	A	40	18.384	75.733	-30.794	1.00	25.57	T2
	ATOM	1457	NZ	LYS	A	40	18.433	76.598	-29.569	1.00	26.12	T2
	ATOM	1458	C	LYS	A	40	15.487	72.342	-33.282	1.00	27.75	T2
25	ATOM	1459	O	LYS	A	40	14.995	72.201	-32.159	1.00	37.35	T2
	ATOM	1460	N	GLU	A	41	16.216	71.414	-33.886	1.00	26.04	T2
	ATOM	1461	CA	GLU	A	41	16.526	70.158	-33.232	1.00	34.07	T2
	ATOM	1462	CB	GLU	A	41	17.561	70.419	-32.142	1.00	33.86	T2
	ATOM	1463	CG	GLU	A	41	18.748	71.206	-32.684	1.00	27.08	T2
30	ATOM	1464	CD	GLU	A	41	19.620	71.801	-31.593	1.00	27.35	T2
	ATOM	1465	OE1	GLU	A	41	19.070	72.495	-30.690	1.00	36.92	T2
	ATOM	1466	OE2	GLU	A	41	20.859	71.583	-31.653	1.00	29.15	T2
	ATOM	1467	C	GLU	A	41	15.284	69.489	-32.658	1.00	37.75	T2
	ATOM	1468	O	GLU	A	41	15.185	69.250	-31.455	1.00	26.04	T2
35	ATOM	1469	N	ASN	A	42	14.332	69.211	-33.538	1.00	32.89	T2
	ATOM	1470	CA	ASN	A	42	13.097	68.547	-33.169	1.00	33.20	T2
	ATOM	1471	CB	ASN	A	42	13.410	67.169	-32.624	1.00	24.23	T2
	ATOM	1472	CG	ASN	A	42	12.461	66.132	-33.138	1.00	29.51	T2
	ATOM	1473	OD1	ASN	A	42	11.840	65.402	-32.367	1.00	32.80	T2
40	ATOM	1474	ND2	ASN	A	42	12.337	66.055	-34.458	1.00	37.44	T2
	ATOM	1475	C	ASN	A	42	12.218	69.290	-32.172	1.00	29.09	T2
	ATOM	1476	O	ASN	A	42	11.370	68.685	-31.508	1.00	26.02	T2
	ATOM	1477	N	LYS	A	43	12.420	70.596	-32.068	1.00	36.16	T2
	ATOM	1478	CA	LYS	A	43	11.634	71.409	-31.164	1.00	29.61	T2
45	ATOM	1479	CB	LYS	A	43	12.459	71.792	-29.943	1.00	32.76	T2
	ATOM	1480	CG	LYS	A	43	12.758	70.641	-29.015	1.00	27.19	T2
	ATOM	1481	CD	LYS	A	43	13.634	71.083	-27.849	1.00	28.51	T2
	ATOM	1482	CE	LYS	A	43	15.051	71.384	-28.313	1.00	32.45	T2
	ATOM	1483	NZ	LYS	A	43	15.948	71.779	-27.188	1.00	30.60	T2
50	ATOM	1484	C	LYS	A	43	11.202	72.661	-31.894	1.00	25.66	T2
	ATOM	1485	O	LYS	A	43	11.773	73.011	-32.923	1.00	34.45	T2
	ATOM	1486	N	ILE	A	44	10.182	73.331	-31.374	1.00	23.47	T2
	ATOM	1487	CA	ILE	A	44	9.719	74.559	-31.992	1.00	28.80	T2
	ATOM	1488	CB	ILE	A	44	8.189	74.673	-31.933	1.00	23.72	T2
55	ATOM	1489	CG2	ILE	A	44	7.744	75.982	-32.571	1.00	24.44	T2
	ATOM	1490	CG1	ILE	A	44	7.558	73.482	-32.657	1.00	24.08	T2
	ATOM	1491	CD1	ILE	A	44	6.050	73.487	-32.655	1.00	29.72	T2
	ATOM	1492	C	ILE	A	44	10.344	75.722	-31.236	1.00	26.82	T2
	ATOM	1493	O	ILE	A	44	10.206	75.833	-30.025	1.00	31.06	T2
60	ATOM	1494	N	LEU	A	45	11.036	76.584	-31.958	1.00	28.90	T2
	ATOM	1495	CA	LEU	A	45	11.696	77.725	-31.349	1.00	30.98	T2
	ATOM	1496	CB	LEU	A	45	13.075	77.905	-31.972	1.00	25.48	T2
	ATOM	1497	CG	LEU	A	45	13.830	79.129	-31.472	1.00	31.47	T2
	ATOM	1498	CD1	LEU	A	45	14.251	78.912	-30.031	1.00	40.26	T2
65	ATOM	1499	CD2	LEU	A	45	15.024	79.364	-32.360	1.00	28.36	T2
	ATOM	1500	C	LEU	A	45	10.911	79.023	-31.491	1.00	35.45	T2

	ATOM	1501	O	LEU	A	45	10.512	79.395	-32.590	1.00	35.01	T2
	ATOM	1502	N	VAL	A	46	10.715	79.720	-30.381	1.00	27.56	T2
	ATOM	1503	CA	VAL	A	46	9.976	80.973	-30.398	1.00	36.42	T2
	ATOM	1504	CB	VAL	A	46	9.332	81.244	-29.037	1.00	34.04	T2
5	ATOM	1505	CG1	VAL	A	46	8.586	82.560	-29.067	1.00	31.29	T2
	ATOM	1506	CG2	VAL	A	46	8.394	80.116	-28.683	1.00	29.51	T2
	ATOM	1507	C	VAL	A	46	10.889	82.136	-30.744	1.00	26.88	T2
	ATOM	1508	O	VAL	A	46	11.853	82.400	-30.037	1.00	32.81	T2
	ATOM	1509	N	LYS	A	47	10.580	82.842	-31.825	1.00	31.78	T2
10	ATOM	1510	CA	LYS	A	47	11.403	83.968	-32.250	1.00	29.40	T2
	ATOM	1511	CB	LYS	A	47	11.650	83.893	-33.759	1.00	21.98	T2
	ATOM	1512	CG	LYS	A	47	12.665	82.856	-34.179	1.00	31.18	T2
	ATOM	1513	CD	LYS	A	47	13.957	83.065	-33.417	1.00	37.34	T2
	ATOM	1514	CE	LYS	A	47	15.115	82.272	-34.001	1.00	22.63	T2
15	ATOM	1515	NZ	LYS	A	47	15.611	82.846	-35.295	1.00	35.67	T2
	ATOM	1516	C	LYS	A	47	10.847	85.345	-31.897	1.00	26.26	T2
	ATOM	1517	O	LYS	A	47	11.527	86.352	-32.062	1.00	31.75	T2
	ATOM	1518	N	GLU	A	48	9.609	85.387	-31.425	1.00	20.26	T2
	ATOM	1519	CA	GLU	A	48	8.963	86.643	-31.050	1.00	31.03	T2
20	ATOM	1520	CB	GLU	A	48	8.068	87.152	-32.167	1.00	34.35	T2
	ATOM	1521	CG	GLU	A	48	8.780	87.650	-33.381	1.00	29.43	T2
	ATOM	1522	CD	GLU	A	48	7.802	88.014	-34.483	1.00	31.13	T2
	ATOM	1523	OE1	GLU	A	48	6.749	88.629	-34.167	1.00	29.86	T2
	ATOM	1524	OE2	GLU	A	48	8.087	87.688	-35.663	1.00	32.81	T2
25	ATOM	1525	C	GLU	A	48	8.087	86.349	-29.862	1.00	22.53	T2
	ATOM	1526	O	GLU	A	48	7.251	85.453	-29.926	1.00	29.08	T2
	ATOM	1527	N	THR	A	49	8.251	87.094	-28.781	1.00	28.09	T2
	ATOM	1528	CA	THR	A	49	7.425	86.823	-27.619	1.00	23.18	T2
	ATOM	1529	CB	THR	A	49	7.938	87.563	-26.377	1.00	31.09	T2
30	ATOM	1530	OG1	THR	A	49	7.307	88.838	-26.298	1.00	28.78	T2
	ATOM	1531	CG2	THR	A	49	9.447	87.761	-26.462	1.00	35.97	T2
	ATOM	1532	C	THR	A	49	5.990	87.244	-27.918	1.00	27.19	T2
	ATOM	1533	O	THR	A	49	5.753	88.153	-28.710	1.00	29.09	T2
	ATOM	1534	N	GLY	A	50	5.039	86.558	-27.292	1.00	30.66	T2
35	ATOM	1535	CA	GLY	A	50	3.639	86.863	-27.500	1.00	29.24	T2
	ATOM	1536	C	GLY	A	50	2.764	85.725	-27.015	1.00	35.96	T2
	ATOM	1537	O	GLY	A	50	3.228	84.860	-26.268	1.00	27.28	T2
	ATOM	1538	N	TYR	A	51	1.497	85.725	-27.431	1.00	36.06	T2
	ATOM	1539	CA	TYR	A	51	0.561	84.676	-27.042	1.00	28.91	T2
40	ATOM	1540	CB	TYR	A	51	-0.786	85.285	-26.650	1.00	33.33	T2
	ATOM	1541	CG	TYR	A	51	-0.739	85.954	-25.305	1.00	30.37	T2
	ATOM	1542	CD1	TYR	A	51	-0.196	87.223	-25.155	1.00	37.07	T2
	ATOM	1543	CE1	TYR	A	51	-0.046	87.796	-23.894	1.00	28.11	T2
	ATOM	1544	CD2	TYR	A	51	-1.142	85.273	-24.163	1.00	33.24	T2
45	ATOM	1545	CE2	TYR	A	51	-0.996	85.830	-22.899	1.00	39.13	T2
	ATOM	1546	CZ	TYR	A	51	-0.443	87.090	-22.766	1.00	36.69	T2
	ATOM	1547	OH	TYR	A	51	-0.247	87.617	-21.500	1.00	33.31	T2
	ATOM	1548	C	TYR	A	51	0.385	83.671	-28.171	1.00	28.76	T2
	ATOM	1549	O	TYR	A	51	0.114	84.043	-29.309	1.00	27.89	T2
50	ATOM	1550	N	PHE	A	52	0.548	82.393	-27.855	1.00	33.01	T2
	ATOM	1551	CA	PHE	A	52	0.426	81.356	-28.867	1.00	30.81	T2
	ATOM	1552	CB	PHE	A	52	1.778	80.677	-29.106	1.00	33.58	T2
	ATOM	1553	CG	PHE	A	52	2.873	81.614	-29.538	1.00	27.55	T2
	ATOM	1554	CD1	PHE	A	52	3.471	82.475	-28.627	1.00	31.29	T2
55	ATOM	1555	CD2	PHE	A	52	3.309	81.625	-30.856	1.00	35.17	T2
	ATOM	1556	CE1	PHE	A	52	4.483	83.327	-29.022	1.00	35.62	T2
	ATOM	1557	CE2	PHE	A	52	4.322	82.475	-31.257	1.00	28.50	T2
	ATOM	1558	CZ	PHE	A	52	4.910	83.328	-30.339	1.00	34.07	T2
	ATOM	1559	C	PHE	A	52	-0.576	80.277	-28.526	1.00	30.68	T2
60	ATOM	1560	O	PHE	A	52	-0.784	79.950	-27.359	1.00	29.36	T2
	ATOM	1561	N	PHE	A	53	-1.195	79.734	-29.568	1.00	33.81	T2
	ATOM	1562	CA	PHE	A	53	-2.142	78.637	-29.435	1.00	30.33	T2
	ATOM	1563	CB	PHE	A	53	-3.269	78.765	-30.455	1.00	37.29	T2
	ATOM	1564	CG	PHE	A	53	-4.164	77.561	-30.522	1.00	38.28	T2
65	ATOM	1565	CD1	PHE	A	53	-4.992	77.232	-29.462	1.00	39.42	T2
	ATOM	1566	CD2	PHE	A	53	-4.167	76.750	-31.645	1.00	35.12	T2

	ATOM	1567	CE1	PHE	A	53	-5.809	76.114	-29.523	1.00	28.29	T2
	ATOM	1568	CE2	PHE	A	53	-4.980	75.633	-31.711	1.00	29.87	T2
	ATOM	1569	CZ	PHE	A	53	-5.800	75.316	-30.648	1.00	34.83	T2
	ATOM	1570	C	PHE	A	53	-1.283	77.438	-29.779	1.00	29.96	T2
5	ATOM	1571	O	PHE	A	53	-0.683	77.401	-30.846	1.00	33.48	T2
	ATOM	1572	N	ILE	A	54	-1.214	76.464	-28.886	1.00	32.00	T2
	ATOM	1573	CA	ILE	A	54	-0.376	75.305	-29.130	1.00	34.73	T2
	ATOM	1574	CB	ILE	A	54	0.726	75.230	-28.073	1.00	26.00	T2
	ATOM	1575	CG2	ILE	A	54	1.726	74.146	-28.437	1.00	31.76	T2
10	ATOM	1576	CG1	ILE	A	54	1.426	76.587	-27.985	1.00	27.88	T2
	ATOM	1577	CD1	ILE	A	54	2.192	76.799	-26.724	1.00	36.19	T2
	ATOM	1578	C	ILE	A	54	-1.183	74.032	-29.104	1.00	32.63	T2
	ATOM	1579	O	ILE	A	54	-1.947	73.808	-28.174	1.00	25.42	T2
	ATOM	1580	N	TYR	A	55	-1.010	73.197	-30.122	1.00	32.35	T2
15	ATOM	1581	CA	TYR	A	55	-1.753	71.947	-30.207	1.00	30.71	T2
	ATOM	1582	CB	TYR	A	55	-2.851	72.072	-31.267	1.00	32.77	T2
	ATOM	1583	CG	TYR	A	55	-2.341	72.439	-32.640	1.00	36.16	T2
	ATOM	1584	CD1	TYR	A	55	-2.078	71.461	-33.592	1.00	32.88	T2
	ATOM	1585	CE1	TYR	A	55	-1.574	71.793	-34.835	1.00	27.02	T2
20	ATOM	1586	CD2	TYR	A	55	-2.089	73.763	-32.971	1.00	25.77	T2
	ATOM	1587	CE2	TYR	A	55	-1.585	74.105	-34.209	1.00	30.61	T2
	ATOM	1588	CZ	TYR	A	55	-1.326	73.117	-35.138	1.00	35.97	T2
	ATOM	1589	OH	TYR	A	55	-0.800	73.457	-36.361	1.00	34.31	T2
	ATOM	1590	C	TYR	A	55	-0.866	70.748	-30.517	1.00	25.96	T2
25	ATOM	1591	O	TYR	A	55	0.311	70.896	-30.811	1.00	31.86	T2
	ATOM	1592	N	GLY	A	56	-1.436	69.555	-30.442	1.00	24.04	T2
	ATOM	1593	CA	GLY	A	56	-0.661	68.371	-30.723	1.00	32.63	T2
	ATOM	1594	C	GLY	A	56	-1.473	67.116	-30.514	1.00	32.59	T2
	ATOM	1595	O	GLY	A	56	-2.249	67.022	-29.567	1.00	29.02	T2
30	ATOM	1596	N	GLN	A	57	-1.292	66.154	-31.416	1.00	28.34	T2
	ATOM	1597	CA	GLN	A	57	-1.988	64.871	-31.353	1.00	31.68	T2
	ATOM	1598	CB	GLN	A	57	-3.113	64.808	-32.398	1.00	26.89	T2
	ATOM	1599	CG	GLN	A	57	-3.774	63.435	-32.499	1.00	29.94	T2
	ATOM	1600	CD	GLN	A	57	-4.971	63.398	-33.432	1.00	27.15	T2
35	ATOM	1601	OE1	GLN	A	57	-6.023	63.971	-33.144	1.00	34.47	T2
	ATOM	1602	NE2	GLN	A	57	-4.814	62.714	-34.560	1.00	23.78	T2
	ATOM	1603	C	GLN	A	57	-1.009	63.726	-31.599	1.00	31.14	T2
	ATOM	1604	O	GLN	A	57	-0.044	63.870	-32.353	1.00	33.24	T2
	ATOM	1605	N	VAL	A	58	-1.267	62.598	-30.945	1.00	29.17	T2
40	ATOM	1606	CA	VAL	A	58	-0.441	61.402	-31.077	1.00	32.28	T2
	ATOM	1607	CB	VAL	A	58	0.507	61.238	-29.866	1.00	31.06	T2
	ATOM	1608	CG1	VAL	A	58	1.149	59.869	-29.883	1.00	33.69	T2
	ATOM	1609	CG2	VAL	A	58	1.569	62.302	-29.895	1.00	39.76	T2
	ATOM	1610	C	VAL	A	58	-1.358	60.184	-31.136	1.00	29.53	T2
45	ATOM	1611	O	VAL	A	58	-2.357	60.140	-30.428	1.00	25.67	T2
	ATOM	1612	N	LEU	A	59	-1.035	59.209	-31.987	1.00	28.55	T2
	ATOM	1613	CA	LEU	A	59	-1.837	57.988	-32.092	1.00	32.42	T2
	ATOM	1614	CB	LEU	A	59	-1.982	57.555	-33.549	1.00	34.54	T2
	ATOM	1615	CG	LEU	A	59	-3.200	56.685	-33.881	1.00	28.26	T2
50	ATOM	1616	CD1	LEU	A	59	-2.968	56.001	-35.215	1.00	35.35	T2
	ATOM	1617	CD2	LEU	A	59	-3.436	55.645	-32.805	1.00	34.46	T2
	ATOM	1618	C	LEU	A	59	-1.120	56.885	-31.307	1.00	34.80	T2
	ATOM	1619	O	LEU	A	59	-0.022	56.454	-31.676	1.00	26.90	T2
	ATOM	1620	N	TYR	A	60	-1.734	56.427	-30.224	1.00	34.28	T2
55	ATOM	1621	CA	TYR	A	60	-1.120	55.384	-29.411	1.00	31.56	T2
	ATOM	1622	CB	TYR	A	60	-1.447	55.611	-27.939	1.00	22.21	T2
	ATOM	1623	CG	TYR	A	60	-0.944	56.930	-27.451	1.00	31.80	T2
	ATOM	1624	CD1	TYR	A	60	-1.820	57.984	-27.219	1.00	29.31	T2
	ATOM	1625	CE1	TYR	A	60	-1.350	59.228	-26.839	1.00	30.24	T2
60	ATOM	1626	CD2	TYR	A	60	0.419	57.143	-27.287	1.00	30.97	T2
	ATOM	1627	CE2	TYR	A	60	0.907	58.370	-26.913	1.00	23.63	T2
	ATOM	1628	CZ	TYR	A	60	0.019	59.421	-26.691	1.00	38.03	T2
	ATOM	1629	OH	TYR	A	60	0.506	60.672	-26.348	1.00	25.07	T2
	ATOM	1630	C	TYR	A	60	-1.529	53.973	-29.800	1.00	29.31	T2
65	ATOM	1631	O	TYR	A	60	-2.713	53.639	-29.866	1.00	30.86	T2
	ATOM	1632	N	THR	A	61	-0.537	53.136	-30.045	1.00	30.76	T2

	ATOM	1633	CA	THR	A	61	-0.800	51.755	-30.415	1.00	26.84	T2
	ATOM	1634	CB	THR	A	61	-0.237	51.449	-31.815	1.00	34.31	T2
	ATOM	1635	OG1	THR	A	61	1.150	51.812	-31.867	1.00	23.48	T2
	ATOM	1636	CG2	THR	A	61	-0.996	52.239	-32.865	1.00	26.43	T2
5	ATOM	1637	C	THR	A	61	-0.145	50.852	-29.383	1.00	29.98	T2
	ATOM	1638	O	THR	A	61	0.034	49.657	-29.598	1.00	36.18	T2
	ATOM	1639	N	ASP	A	62	0.211	51.453	-28.256	1.00	30.32	T2
	ATOM	1640	CA	ASP	A	62	0.843	50.750	-27.150	1.00	34.04	T2
	ATOM	1641	CB	ASP	A	62	1.684	51.755	-26.358	1.00	21.40	T2
10	ATOM	1642	CG	ASP	A	62	2.621	51.094	-25.368	1.00	36.43	T2
	ATOM	1643	OD1	ASP	A	62	3.831	51.457	-25.372	1.00	31.95	T2
	ATOM	1644	OD2	ASP	A	62	2.146	50.228	-24.584	1.00	27.87	T2
	ATOM	1645	C	ASP	A	62	-0.286	50.184	-26.292	1.00	27.39	T2
	ATOM	1646	O	ASP	A	62	-1.361	50.770	-26.217	1.00	33.83	T2
15	ATOM	1647	N	LYS	A	63	-0.067	49.050	-25.646	1.00	24.01	T2
	ATOM	1648	CA	LYS	A	63	-1.137	48.477	-24.824	1.00	32.86	T2
	ATOM	1649	CB	LYS	A	63	-1.353	47.007	-25.191	1.00	34.03	T2
	ATOM	1650	CG	LYS	A	63	-0.119	46.135	-24.979	1.00	31.83	T2
	ATOM	1651	CD	LYS	A	63	-0.407	44.672	-25.331	1.00	25.10	T2
20	ATOM	1652	CE	LYS	A	63	-0.826	44.505	-26.807	1.00	37.90	T2
	ATOM	1653	NZ	LYS	A	63	-1.150	43.078	-27.172	1.00	29.56	T2
	ATOM	1654	C	LYS	A	63	-0.914	48.587	-23.312	1.00	26.42	T2
	ATOM	1655	O	LYS	A	63	-1.510	47.833	-22.534	1.00	30.96	T2
	ATOM	1656	N	THR	A	64	-0.088	49.533	-22.885	1.00	31.94	T2
25	ATOM	1657	CA	THR	A	64	0.170	49.647	-21.460	1.00	25.58	T2
	ATOM	1658	CB	THR	A	64	1.607	50.185	-21.190	1.00	36.84	T2
	ATOM	1659	OG1	THR	A	64	1.841	51.384	-21.940	1.00	34.01	T2
	ATOM	1660	CG2	THR	A	64	2.639	49.129	-21.588	1.00	31.83	T2
	ATOM	1661	C	THR	A	64	-0.844	50.437	-20.633	1.00	29.65	T2
30	ATOM	1662	O	THR	A	64	-0.499	51.431	-19.990	1.00	32.84	T2
	ATOM	1663	N	TYR	A	65	-2.093	49.976	-20.654	1.00	24.57	T2
	ATOM	1664	CA	TYR	A	65	-3.199	50.571	-19.884	1.00	32.07	T2
	ATOM	1665	CB	TYR	A	65	-3.120	50.083	-18.429	1.00	31.30	T2
	ATOM	1666	CG	TYR	A	65	-2.562	51.090	-17.454	1.00	32.12	T2
35	ATOM	1667	CD1	TYR	A	65	-3.408	51.963	-16.756	1.00	26.54	T2
	ATOM	1668	CE1	TYR	A	65	-2.891	52.909	-15.845	1.00	25.19	T2
	ATOM	1669	CD2	TYR	A	65	-1.189	51.181	-17.227	1.00	32.08	T2
	ATOM	1670	CE2	TYR	A	65	-0.657	52.121	-16.325	1.00	33.18	T2
	ATOM	1671	CZ	TYR	A	65	-1.510	52.978	-15.638	1.00	27.84	T2
40	ATOM	1672	OH	TYR	A	65	-0.973	53.897	-14.757	1.00	29.92	T2
	ATOM	1673	C	TYR	A	65	-3.403	52.098	-19.886	1.00	33.02	T2
	ATOM	1674	O	TYR	A	65	-4.518	52.568	-19.632	1.00	30.03	T2
	ATOM	1675	N	ALA	A	66	-2.355	52.871	-20.151	1.00	32.82	T2
	ATOM	1676	CA	ALA	A	66	-2.501	54.314	-20.159	1.00	31.49	T2
45	ATOM	1677	CB	ALA	A	66	-2.592	54.832	-18.729	1.00	33.63	T2
	ATOM	1678	C	ALA	A	66	-1.352	54.990	-20.892	1.00	28.59	T2
	ATOM	1679	O	ALA	A	66	-0.191	54.870	-20.493	1.00	28.70	T2
	ATOM	1680	N	MET	A	67	-1.678	55.694	-21.972	1.00	26.02	T2
	ATOM	1681	CA	MET	A	67	-0.672	56.412	-22.736	1.00	38.12	T2
50	ATOM	1682	CB	MET	A	67	-0.608	55.869	-24.160	1.00	31.84	T2
	ATOM	1683	CG	MET	A	67	-0.029	54.469	-24.249	1.00	33.75	T2
	ATOM	1684	SD	MET	A	67	1.657	54.413	-23.588	1.00	30.42	T2
	ATOM	1685	CE	MET	A	67	2.596	55.054	-25.000	1.00	32.56	T2
	ATOM	1686	C	MET	A	67	-1.020	57.894	-22.759	1.00	29.78	T2
55	ATOM	1687	O	MET	A	67	-2.119	58.286	-22.360	1.00	32.68	T2
	ATOM	1688	N	GLY	A	68	-0.083	58.716	-23.218	1.00	33.35	T2
	ATOM	1689	CA	GLY	A	68	-0.331	60.144	-23.282	1.00	30.46	T2
	ATOM	1690	C	GLY	A	68	0.927	60.933	-23.570	1.00	34.96	T2
	ATOM	1691	O	GLY	A	68	2.031	60.380	-23.550	1.00	31.69	T2
60	ATOM	1692	N	HIS	A	69	0.770	62.223	-23.859	1.00	31.90	T2
	ATOM	1693	CA	HIS	A	69	1.929	63.074	-24.130	1.00	23.28	T2
	ATOM	1694	CB	HIS	A	69	2.124	63.287	-25.639	1.00	31.42	T2
	ATOM	1695	CG	HIS	A	69	0.915	63.812	-26.346	1.00	33.89	T2
	ATOM	1696	CD2	HIS	A	69	0.627	65.047	-26.818	1.00	34.46	T2
65	ATOM	1697	ND1	HIS	A	69	-0.163	63.017	-26.672	1.00	36.73	T2
	ATOM	1698	CE1	HIS	A	69	-1.059	63.739	-27.319	1.00	31.36	T2

	ATOM	1699	NE2	HIS	A	69	-0.604	64.974	-27.421	1.00	33.49	T2
	ATOM	1700	C	HIS	A	69	1.855	64.424	-23.422	1.00	27.39	T2
	ATOM	1701	O	HIS	A	69	0.808	64.814	-22.892	1.00	34.50	T2
	ATOM	1702	N	LEU	A	70	2.985	65.123	-23.420	1.00	35.46	T2
5	ATOM	1703	CA	LEU	A	70	3.085	66.417	-22.774	1.00	26.54	T2
	ATOM	1704	CB	LEU	A	70	4.099	66.361	-21.636	1.00	34.22	T2
	ATOM	1705	CG	LEU	A	70	4.130	65.133	-20.742	1.00	28.15	T2
	ATOM	1706	CD1	LEU	A	70	5.333	65.225	-19.839	1.00	25.52	T2
	ATOM	1707	CD2	LEU	A	70	2.862	65.036	-19.933	1.00	26.36	T2
10	ATOM	1708	C	LEU	A	70	3.569	67.465	-23.761	1.00	32.24	T2
	ATOM	1709	O	LEU	A	70	4.381	67.169	-24.635	1.00	32.33	T2
	ATOM	1710	N	ILE	A	71	3.066	68.686	-23.623	1.00	36.30	T2
	ATOM	1711	CA	ILE	A	71	3.524	69.777	-24.460	1.00	33.79	T2
	ATOM	1712	CB	ILE	A	71	2.360	70.549	-25.070	1.00	28.07	T2
15	ATOM	1713	CG2	ILE	A	71	2.854	71.845	-25.667	1.00	30.24	T2
	ATOM	1714	CG1	ILE	A	71	1.697	69.700	-26.153	1.00	29.51	T2
	ATOM	1715	CD1	ILE	A	71	0.504	70.360	-26.800	1.00	22.52	T2
	ATOM	1716	C	ILE	A	71	4.287	70.650	-23.475	1.00	38.15	T2
	ATOM	1717	O	ILE	A	71	3.694	71.278	-22.601	1.00	28.96	T2
20	ATOM	1718	N	GLN	A	72	5.610	70.679	-23.604	1.00	26.25	T2
	ATOM	1719	CA	GLN	A	72	6.426	71.439	-22.666	1.00	31.83	T2
	ATOM	1720	CB	GLN	A	72	7.480	70.516	-22.073	1.00	32.21	T2
	ATOM	1721	CG	GLN	A	72	6.898	69.191	-21.651	1.00	32.65	T2
	ATOM	1722	CD	GLN	A	72	7.881	68.358	-20.877	1.00	37.31	T2
25	ATOM	1723	OE1	GLN	A	72	8.994	68.088	-21.347	1.00	27.06	T2
	ATOM	1724	NE2	GLN	A	72	7.482	67.937	-19.677	1.00	26.87	T2
	ATOM	1725	C	GLN	A	72	7.089	72.697	-23.194	1.00	33.67	T2
	ATOM	1726	O	GLN	A	72	7.316	72.859	-24.393	1.00	29.63	T2
	ATOM	1727	N	ARG	A	73	7.411	73.581	-22.259	1.00	26.18	T2
30	ATOM	1728	CA	ARG	A	73	8.048	74.849	-22.561	1.00	31.30	T2
	ATOM	1729	CB	ARG	A	73	7.115	75.978	-22.147	1.00	36.20	T2
	ATOM	1730	CG	ARG	A	73	7.669	77.342	-22.398	1.00	26.55	T2
	ATOM	1731	CD	ARG	A	73	7.037	78.352	-21.491	1.00	33.43	T2
	ATOM	1732	NE	ARG	A	73	7.597	79.678	-21.714	1.00	29.92	T2
35	ATOM	1733	CZ	ARG	A	73	7.439	80.699	-20.883	1.00	27.09	T2
	ATOM	1734	NH1	ARG	A	73	6.735	80.540	-19.766	1.00	28.57	T2
	ATOM	1735	NH2	ARG	A	73	7.981	81.874	-21.173	1.00	34.61	T2
	ATOM	1736	C	ARG	A	73	9.379	74.996	-21.810	1.00	26.94	T2
	ATOM	1737	O	ARG	A	73	9.449	74.731	-20.603	1.00	27.69	T2
40	ATOM	1738	N	LYS	A	74	10.429	75.411	-22.524	1.00	28.38	T2
	ATOM	1739	CA	LYS	A	74	11.744	75.623	-21.915	1.00	27.94	T2
	ATOM	1740	CB	LYS	A	74	12.850	75.001	-22.763	1.00	30.54	T2
	ATOM	1741	CG	LYS	A	74	12.829	73.480	-22.808	1.00	30.10	T2
	ATOM	1742	CD	LYS	A	74	13.918	72.910	-23.739	1.00	37.12	T2
45	ATOM	1743	CE	LYS	A	74	13.866	71.369	-23.791	1.00	30.02	T2
	ATOM	1744	NZ	LYS	A	74	14.929	70.763	-24.663	1.00	35.90	T2
	ATOM	1745	C	LYS	A	74	11.988	77.112	-21.818	1.00	37.44	T2
	ATOM	1746	O	LYS	A	74	12.356	77.744	-22.807	1.00	32.05	T2
	ATOM	1747	N	LYS	A	75	11.782	77.668	-20.628	1.00	29.69	T2
50	ATOM	1748	CA	LYS	A	75	11.967	79.098	-20.400	1.00	27.40	T2
	ATOM	1749	CB	LYS	A	75	11.648	79.454	-18.949	1.00	35.38	T2
	ATOM	1750	CG	LYS	A	75	10.246	79.155	-18.492	1.00	41.11	T2
	ATOM	1751	CD	LYS	A	75	10.084	79.573	-17.039	1.00	35.02	T2
	ATOM	1752	CE	LYS	A	75	8.682	79.266	-16.520	1.00	33.10	T2
55	ATOM	1753	NZ	LYS	A	75	8.494	79.546	-15.049	1.00	31.50	T2
	ATOM	1754	C	LYS	A	75	13.397	79.538	-20.696	1.00	30.81	T2
	ATOM	1755	O	LYS	A	75	14.341	78.863	-20.302	1.00	34.73	T2
	ATOM	1756	N	VAL	A	76	13.554	80.670	-21.378	1.00	26.11	T2
	ATOM	1757	CA	VAL	A	76	14.882	81.192	-21.687	1.00	31.99	T2
60	ATOM	1758	CB	VAL	A	76	14.862	82.286	-22.747	1.00	30.54	T2
	ATOM	1759	CG1	VAL	A	76	16.250	82.478	-23.298	1.00	33.21	T2
	ATOM	1760	CG2	VAL	A	76	13.891	81.948	-23.821	1.00	34.73	T2
	ATOM	1761	C	VAL	A	76	15.388	81.867	-20.436	1.00	39.20	T2
	ATOM	1762	O	VAL	A	76	16.557	81.751	-20.080	1.00	35.62	T2
65	ATOM	1763	N	HIS	A	77	14.486	82.591	-19.784	1.00	28.61	T2
	ATOM	1764	CA	HIS	A	77	14.802	83.312	-18.568	1.00	29.79	T2

	ATOM	1765	CB	HIS	A	77	14.228	84.718	-18.646	1.00	36.12	T2
	ATOM	1766	CG	HIS	A	77	14.722	85.500	-19.821	1.00	32.56	T2
	ATOM	1767	CD2	HIS	A	77	14.220	86.593	-20.443	1.00	31.03	T2
	ATOM	1768	ND1	HIS	A	77	15.915	85.215	-20.451	1.00	26.25	T2
5	ATOM	1769	CE1	HIS	A	77	16.130	86.101	-21.407	1.00	26.58	T2
	ATOM	1770	NE2	HIS	A	77	15.117	86.949	-21.421	1.00	36.58	T2
	ATOM	1771	C	HIS	A	77	14.225	82.557	-17.389	1.00	27.12	T2
	ATOM	1772	O	HIS	A	77	13.175	81.935	-17.505	1.00	35.28	T2
	ATOM	1773	N	VAL	A	78	14.901	82.620	-16.246	1.00	28.70	T2
10	ATOM	1774	CA	VAL	A	78	14.430	81.875	-15.099	1.00	32.50	T2
	ATOM	1775	CB	VAL	A	78	15.374	80.682	-14.849	1.00	27.69	T2
	ATOM	1776	CG1	VAL	A	78	14.823	79.797	-13.761	1.00	26.67	T2
	ATOM	1777	CG2	VAL	A	78	15.503	79.864	-16.117	1.00	30.75	T2
	ATOM	1778	C	VAL	A	78	14.117	82.597	-13.773	1.00	28.22	T2
15	ATOM	1779	O	VAL	A	78	12.958	82.635	-13.362	1.00	30.12	T2
	ATOM	1780	N	PHE	A	79	15.098	83.165	-13.088	1.00	26.22	T2
	ATOM	1781	CA	PHE	A	79	14.805	83.835	-11.800	1.00	29.48	T2
	ATOM	1782	CB	PHE	A	79	13.546	84.712	-11.869	1.00	28.07	T2
	ATOM	1783	CG	PHE	A	79	13.542	85.693	-12.987	1.00	33.87	T2
20	ATOM	1784	CD1	PHE	A	79	12.826	85.433	-14.143	1.00	19.16	T2
	ATOM	1785	CD2	PHE	A	79	14.259	86.876	-12.894	1.00	31.70	T2
	ATOM	1786	CE1	PHE	A	79	12.822	86.335	-15.197	1.00	24.00	T2
	ATOM	1787	CE2	PHE	A	79	14.264	87.790	-13.942	1.00	36.64	T2
	ATOM	1788	CZ	PHE	A	79	13.544	87.518	-15.097	1.00	27.65	T2
25	ATOM	1789	C	PHE	A	79	14.596	82.896	-10.604	1.00	36.91	T2
	ATOM	1790	O	PHE	A	79	13.714	82.043	-10.619	1.00	30.13	T2
	ATOM	1791	N	GLY	A	80	15.397	83.092	-9.561	1.00	31.63	T2
	ATOM	1792	CA	GLY	A	80	15.286	82.299	-8.347	1.00	36.14	T2
	ATOM	1793	C	GLY	A	80	15.175	80.797	-8.505	1.00	31.94	T2
30	ATOM	1794	O	GLY	A	80	15.969	80.171	-9.218	1.00	34.22	T2
	ATOM	1795	N	ASP	A	81	14.180	80.220	-7.832	1.00	27.88	T2
	ATOM	1796	CA	ASP	A	81	13.952	78.780	-7.870	1.00	28.81	T2
	ATOM	1797	CB	ASP	A	81	13.610	78.272	-6.468	1.00	31.49	T2
	ATOM	1798	CG	ASP	A	81	12.290	78.811	-5.948	1.00	32.48	T2
35	ATOM	1799	OD1	ASP	A	81	11.785	79.803	-6.504	1.00	25.07	T2
	ATOM	1800	OD2	ASP	A	81	11.759	78.251	-4.965	1.00	33.89	T2
	ATOM	1801	C	ASP	A	81	12.883	78.322	-8.864	1.00	26.40	T2
	ATOM	1802	O	ASP	A	81	12.210	77.311	-8.642	1.00	33.00	T2
	ATOM	1803	N	GLU	A	82	12.722	79.061	-9.957	1.00	31.32	T2
40	ATOM	1804	CA	GLU	A	82	11.753	78.679	-10.971	1.00	32.41	T2
	ATOM	1805	CB	GLU	A	82	11.601	79.755	-12.035	1.00	34.15	T2
	ATOM	1806	CG	GLU	A	82	10.728	80.919	-11.705	1.00	29.63	T2
	ATOM	1807	CD	GLU	A	82	10.136	81.504	-12.971	1.00	37.18	T2
	ATOM	1808	OE1	GLU	A	82	10.876	81.631	-13.970	1.00	28.61	T2
45	ATOM	1809	OE2	GLU	A	82	8.932	81.832	-12.985	1.00	35.11	T2
	ATOM	1810	C	GLU	A	82	12.307	77.463	-11.675	1.00	26.47	T2
	ATOM	1811	O	GLU	A	82	13.523	77.321	-11.768	1.00	36.03	T2
	ATOM	1812	N	LEU	A	83	11.437	76.583	-12.163	1.00	30.79	T2
	ATOM	1813	CA	LEU	A	83	11.915	75.434	-12.923	1.00	30.23	T2
50	ATOM	1814	CB	LEU	A	83	10.994	74.223	-12.772	1.00	30.21	T2
	ATOM	1815	CG	LEU	A	83	10.952	73.546	-11.407	1.00	32.66	T2
	ATOM	1816	CD1	LEU	A	83	10.396	74.519	-10.366	1.00	30.71	T2
	ATOM	1817	CD2	LEU	A	83	10.081	72.313	-11.501	1.00	24.96	T2
	ATOM	1818	C	LEU	A	83	11.845	75.964	-14.340	1.00	37.52	T2
55	ATOM	1819	O	LEU	A	83	10.832	76.542	-14.734	1.00	28.79	T2
	ATOM	1820	N	SER	A	84	12.917	75.793	-15.098	1.00	30.08	T2
	ATOM	1821	CA	SER	A	84	12.941	76.299	-16.462	1.00	31.43	T2
	ATOM	1822	CB	SER	A	84	14.384	76.476	-16.930	1.00	35.65	T2
	ATOM	1823	OG	SER	A	84	15.141	75.318	-16.646	1.00	29.58	T2
60	ATOM	1824	C	SER	A	84	12.173	75.436	-17.450	1.00	30.25	T2
	ATOM	1825	O	SER	A	84	12.046	75.788	-18.622	1.00	28.53	T2
	ATOM	1826	N	LEU	A	85	11.655	74.309	-16.975	1.00	32.88	T2
	ATOM	1827	CA	LEU	A	85	10.887	73.413	-17.829	1.00	32.05	T2
	ATOM	1828	CB	LEU	A	85	11.538	72.020	-17.857	1.00	28.33	T2
65	ATOM	1829	CG	LEU	A	85	11.171	71.013	-18.957	1.00	37.45	T2
	ATOM	1830	CD1	LEU	A	85	9.760	70.493	-18.762	1.00	30.73	T2

	ATOM	1831	CD2	LEU	A	85	11.322	71.670	-20.317	1.00	32.47	T2
	ATOM	1832	C	LEU	A	85	9.466	73.327	-17.282	1.00	33.45	T2
	ATOM	1833	O	LEU	A	85	9.224	72.719	-16.239	1.00	36.80	T2
	ATOM	1834	N	VAL	A	86	8.530	73.962	-17.979	1.00	30.04	T2
5	ATOM	1835	CA	VAL	A	86	7.129	73.943	-17.564	1.00	27.85	T2
	ATOM	1836	CB	VAL	A	86	6.497	75.342	-17.582	1.00	36.12	T2
	ATOM	1837	CG1	VAL	A	86	5.026	75.234	-17.240	1.00	29.68	T2
	ATOM	1838	CG2	VAL	A	86	7.212	76.256	-16.602	1.00	32.34	T2
	ATOM	1839	C	VAL	A	86	6.349	73.094	-18.538	1.00	30.01	T2
10	ATOM	1840	O	VAL	A	86	6.573	73.159	-19.746	1.00	30.77	T2
	ATOM	1841	N	THR	A	87	5.438	72.288	-18.022	1.00	30.66	T2
	ATOM	1842	CA	THR	A	87	4.648	71.465	-18.907	1.00	36.09	T2
	ATOM	1843	CB	THR	A	87	4.645	69.973	-18.473	1.00	29.96	T2
	ATOM	1844	OG1	THR	A	87	3.317	69.575	-18.134	1.00	31.25	T2
15	ATOM	1845	CG2	THR	A	87	5.574	69.750	-17.289	1.00	32.89	T2
	ATOM	1846	C	THR	A	87	3.238	72.027	-18.941	1.00	24.00	T2
	ATOM	1847	O	THR	A	87	2.561	72.109	-17.921	1.00	30.89	T2
	ATOM	1848	N	LEU	A	88	2.827	72.452	-20.133	1.00	31.55	T2
	ATOM	1849	CA	LEU	A	88	1.502	73.016	-20.374	1.00	28.10	T2
20	ATOM	1850	CB	LEU	A	88	1.594	74.103	-21.447	1.00	33.98	T2
	ATOM	1851	CG	LEU	A	88	2.509	75.331	-21.367	1.00	32.82	T2
	ATOM	1852	CD1	LEU	A	88	3.737	75.049	-20.572	1.00	34.96	T2
	ATOM	1853	CD2	LEU	A	88	2.890	75.738	-22.762	1.00	36.74	T2
	ATOM	1854	C	LEU	A	88	0.627	71.895	-20.931	1.00	27.39	T2
25	ATOM	1855	O	LEU	A	88	1.110	71.018	-21.655	1.00	31.06	T2
	ATOM	1856	N	PHE	A	89	-0.648	71.877	-20.587	1.00	34.62	T2
	ATOM	1857	CA	PHE	A	89	-1.529	70.871	-21.192	1.00	34.29	T2
	ATOM	1858	CB	PHE	A	89	-1.488	71.049	-22.717	1.00	27.10	T2
	ATOM	1859	CG	PHE	A	89	-1.525	72.489	-23.133	1.00	26.91	T2
30	ATOM	1860	CD1	PHE	A	89	-0.822	72.928	-24.240	1.00	32.82	T2
	ATOM	1861	CD2	PHE	A	89	-2.199	73.436	-22.339	1.00	34.29	T2
	ATOM	1862	CE1	PHE	A	89	-0.778	74.295	-24.545	1.00	35.66	T2
	ATOM	1863	CE2	PHE	A	89	-2.162	74.799	-22.633	1.00	33.97	T2
	ATOM	1864	CZ	PHE	A	89	-1.450	75.232	-23.733	1.00	27.18	T2
35	ATOM	1865	C	PHE	A	89	-1.349	69.398	-20.832	1.00	37.07	T2
	ATOM	1866	O	PHE	A	89	-1.746	68.996	-19.746	1.00	23.82	T2
	ATOM	1867	N	ARG	A	90	-0.794	68.577	-21.725	1.00	37.10	T2
	ATOM	1868	CA	ARG	A	90	-0.659	67.144	-21.409	1.00	35.07	T2
	ATOM	1869	CB	ARG	A	90	0.075	66.971	-20.076	1.00	36.45	T2
40	ATOM	1870	CG	ARG	A	90	-0.298	65.699	-19.325	1.00	31.81	T2
	ATOM	1871	CD	ARG	A	90	0.125	65.772	-17.880	1.00	23.60	T2
	ATOM	1872	NE	ARG	A	90	-0.437	64.658	-17.133	1.00	31.14	T2
	ATOM	1873	CZ	ARG	A	90	-0.868	64.738	-15.875	1.00	27.61	T2
	ATOM	1874	NH1	ARG	A	90	-0.798	65.894	-15.212	1.00	30.48	T2
45	ATOM	1875	NH2	ARG	A	90	-1.388	63.664	-15.283	1.00	38.53	T2
	ATOM	1876	C	ARG	A	90	-2.007	66.372	-21.326	1.00	31.12	T2
	ATOM	1877	O	ARG	A	90	-2.927	66.766	-20.599	1.00	25.51	T2
	ATOM	1878	N	CYS	A	91	-2.101	65.254	-22.045	1.00	30.24	T2
	ATOM	1879	CA	CYS	A	91	-3.318	64.433	-22.043	1.00	32.91	T2
50	ATOM	1880	CB	CYS	A	91	-4.036	64.573	-23.379	1.00	28.78	T2
	ATOM	1881	SG	CYS	A	91	-2.971	64.169	-24.783	1.00	24.09	T2
	ATOM	1882	C	CYS	A	91	-3.020	62.949	-21.770	1.00	24.03	T2
	ATOM	1883	O	CYS	A	91	-1.865	62.516	-21.833	1.00	34.63	T2
	ATOM	1884	N	ILE	A	92	-4.065	62.178	-21.473	1.00	33.98	T2
55	ATOM	1885	CA	ILE	A	92	-3.927	60.749	-21.154	1.00	30.49	T2
	ATOM	1886	CB	ILE	A	92	-4.025	60.513	-19.639	1.00	36.69	T2
	ATOM	1887	CG2	ILE	A	92	-3.812	59.058	-19.322	1.00	27.60	T2
	ATOM	1888	CG1	ILE	A	92	-2.977	61.349	-18.910	1.00	33.96	T2
	ATOM	1889	CD1	ILE	A	92	-3.208	61.434	-17.400	1.00	30.90	T2
60	ATOM	1890	C	ILE	A	92	-5.049	59.947	-21.794	1.00	34.37	T2
	ATOM	1891	O	ILE	A	92	-6.119	60.474	-22.040	1.00	30.94	T2
	ATOM	1892	N	GLN	A	93	-4.806	58.670	-22.054	1.00	36.11	T2
	ATOM	1893	CA	GLN	A	93	-5.820	57.803	-22.649	1.00	28.85	T2
	ATOM	1894	CB	GLN	A	93	-5.776	57.886	-24.174	1.00	31.75	T2
65	ATOM	1895	CG	GLN	A	93	-6.597	59.018	-24.774	1.00	27.89	T2
	ATOM	1896	CD	GLN	A	93	-7.976	58.566	-25.244	1.00	28.47	T2

	ATOM	1897	OE1	GLN	A	93	-8.887	58.328	-24.443	1.00	33.71	T2
	ATOM	1898	NE2	GLN	A	93	-8.130	58.439	-26.558	1.00	34.61	T2
	ATOM	1899	C	GLN	A	93	-5.623	56.355	-22.232	1.00	39.19	T2
	ATOM	1900	O	GLN	A	93	-4.565	55.762	-22.490	1.00	21.88	T2
5	ATOM	1901	N	ASN	A	94	-6.633	55.787	-21.578	1.00	33.56	T2
	ATOM	1902	CA	ASN	A	94	-6.544	54.398	-21.156	1.00	33.39	T2
	ATOM	1903	CB	ASN	A	94	-7.807	53.976	-20.401	1.00	34.84	T2
	ATOM	1904	CG	ASN	A	94	-7.791	54.414	-18.945	1.00	34.45	T2
	ATOM	1905	OD1	ASN	A	94	-6.849	54.118	-18.217	1.00	29.04	T2
10	ATOM	1906	ND2	ASN	A	94	-8.836	55.112	-18.511	1.00	21.87	T2
	ATOM	1907	C	ASN	A	94	-6.397	53.582	-22.428	1.00	34.09	T2
	ATOM	1908	O	ASN	A	94	-6.946	53.949	-23.465	1.00	25.10	T2
	ATOM	1909	N	MET	A	95	-5.641	52.493	-22.357	1.00	36.93	T2
	ATOM	1910	CA	MET	A	95	-5.437	51.640	-23.518	1.00	33.51	T2
15	ATOM	1911	CB	MET	A	95	-3.943	51.465	-23.801	1.00	35.39	T2
	ATOM	1912	CG	MET	A	95	-3.218	52.749	-24.126	1.00	26.89	T2
	ATOM	1913	SD	MET	A	95	-4.035	53.696	-25.450	1.00	39.72	T2
	ATOM	1914	CB	MET	A	95	-3.585	52.772	-26.922	1.00	32.47	T2
	ATOM	1915	C	MET	A	95	-6.059	50.266	-23.298	1.00	25.58	T2
20	ATOM	1916	O	MET	A	95	-6.168	49.798	-22.159	1.00	36.08	T2
	ATOM	1917	N	PRO	A	96	-6.490	49.605	-24.392	1.00	32.64	T2
	ATOM	1918	CD	PRO	A	96	-6.541	50.113	-25.772	1.00	33.07	T2
	ATOM	1919	CA	PRO	A	96	-7.094	48.272	-24.323	1.00	25.64	T2
	ATOM	1920	CB	PRO	A	96	-7.749	48.094	-25.697	1.00	37.67	T2
25	ATOM	1921	CG	PRO	A	96	-7.819	49.482	-26.264	1.00	27.17	T2
	ATOM	1922	C	PRO	A	96	-5.921	47.310	-24.169	1.00	32.60	T2
	ATOM	1923	O	PRO	A	96	-4.764	47.727	-23.964	1.00	28.36	T2
	ATOM	1924	N	GLU	A	97	-6.201	46.023	-24.305	1.00	31.93	T2
	ATOM	1925	CA	GLU	A	97	-5.142	45.040	-24.170	1.00	33.08	T2
30	ATOM	1926	CB	GLU	A	97	-5.524	44.035	-23.102	1.00	35.77	T2
	ATOM	1927	CG	GLU	A	97	-4.358	43.224	-22.632	1.00	20.63	T2
	ATOM	1928	CD	GLU	A	97	-4.361	43.084	-21.117	1.00	24.69	T2
	ATOM	1929	OE1	GLU	A	97	-5.351	42.530	-20.569	1.00	37.09	T2
	ATOM	1930	OE2	GLU	A	97	-3.377	43.539	-20.470	1.00	32.98	T2
35	ATOM	1931	C	GLU	A	97	-4.916	44.331	-25.493	1.00	27.03	T2
	ATOM	1932	O	GLU	A	97	-3.839	43.794	-25.755	1.00	29.38	T2
	ATOM	1933	N	THR	A	98	-5.938	44.365	-26.335	1.00	31.23	T2
	ATOM	1934	CA	THR	A	98	-5.879	43.697	-27.619	1.00	26.16	T2
	ATOM	1935	CB	THR	A	98	-7.243	43.109	-27.965	1.00	36.85	T2
40	ATOM	1936	OG1	THR	A	98	-8.190	44.183	-28.110	1.00	30.10	T2
	ATOM	1937	CG2	THR	A	98	-7.696	42.150	-26.859	1.00	31.76	T2
	ATOM	1938	C	THR	A	98	-5.419	44.544	-28.799	1.00	30.83	T2
	ATOM	1939	O	THR	A	98	-4.299	44.369	-29.288	1.00	31.63	T2
	ATOM	1940	N	LEU	A	99	-6.263	45.454	-29.269	1.00	31.94	T2
45	ATOM	1941	CA	LEU	A	99	-5.864	46.235	-30.427	1.00	29.69	T2
	ATOM	1942	CB	LEU	A	99	-6.851	45.980	-31.560	1.00	33.59	T2
	ATOM	1943	CG	LEU	A	99	-6.848	44.510	-31.988	1.00	32.92	T2
	ATOM	1944	CD1	LEU	A	99	-7.948	44.251	-33.010	1.00	24.56	T2
	ATOM	1945	CD2	LEU	A	99	-5.488	44.169	-32.571	1.00	27.36	T2
50	ATOM	1946	C	LEU	A	99	-5.704	47.727	-30.188	1.00	27.26	T2
	ATOM	1947	O	LEU	A	99	-6.513	48.537	-30.664	1.00	23.49	T2
	ATOM	1948	N	PRO	A	100	-4.636	48.114	-29.465	1.00	32.93	T2
	ATOM	1949	CD	PRO	A	100	-3.562	47.239	-28.964	1.00	21.55	T2
	ATOM	1950	CA	PRO	A	100	-4.338	49.511	-29.143	1.00	33.22	T2
55	ATOM	1951	CB	PRO	A	100	-2.897	49.443	-28.640	1.00	29.72	T2
	ATOM	1952	CG	PRO	A	100	-2.870	48.130	-27.947	1.00	30.17	T2
	ATOM	1953	C	PRO	A	100	-4.496	50.457	-30.334	1.00	30.41	T2
	ATOM	1954	O	PRO	A	100	-3.857	50.283	-31.375	1.00	29.63	T2
	ATOM	1955	N	ASN	A	101	-5.351	51.456	-30.156	1.00	34.13	T2
60	ATOM	1956	CA	ASN	A	101	-5.613	52.440	-31.184	1.00	35.63	T2
	ATOM	1957	CB	ASN	A	101	-6.504	51.852	-32.256	1.00	31.44	T2
	ATOM	1958	CG	ASN	A	101	-5.727	51.153	-33.310	1.00	23.78	T2
	ATOM	1959	OD1	ASN	A	101	-5.003	51.792	-34.086	1.00	27.95	T2
	ATOM	1960	ND2	ASN	A	101	-5.849	49.825	-33.356	1.00	30.50	T2
65	ATOM	1961	C	ASN	A	101	-6.331	53.619	-30.589	1.00	33.79	T2
	ATOM	1962	O	ASN	A	101	-7.565	53.699	-30.695	1.00	25.62	T2

	ATOM	1963	N	ASN	A	102	-5.602	54.550	-29.974	1.00	27.62	T2
	ATOM	1964	CA	ASN	A	102	-6.325	55.660	-29.404	1.00	30.41	T2
	ATOM	1965	CB	ASN	A	102	-6.237	55.616	-27.884	1.00	34.80	T2
	ATOM	1966	CG	ASN	A	102	-7.296	54.696	-27.291	1.00	30.89	T2
5	ATOM	1967	OD1	ASN	A	102	-8.469	54.745	-27.683	1.00	32.91	T2
	ATOM	1968	ND2	ASN	A	102	-6.890	53.850	-26.355	1.00	36.00	T2
	ATOM	1969	C	ASN	A	102	-6.160	57.075	-29.917	1.00	34.18	T2
	ATOM	1970	O	ASN	A	102	-7.179	57.735	-30.146	1.00	34.28	T2
	ATOM	1971	N	SER	A	103	-4.951	57.579	-30.129	1.00	32.50	T2
10	ATOM	1972	CA	SER	A	103	-4.888	58.962	-30.631	1.00	33.73	T2
	ATOM	1973	CB	SER	A	103	-5.582	59.061	-32.007	1.00	28.14	T2
	ATOM	1974	OG	SER	A	103	-6.482	60.163	-32.071	1.00	24.50	T2
	ATOM	1975	C	SER	A	103	-5.551	59.949	-29.642	1.00	34.64	T2
	ATOM	1976	O	SER	A	103	-6.681	59.760	-29.176	1.00	31.12	T2
15	ATOM	1977	N	CYS	A	104	-4.846	61.021	-29.328	1.00	24.55	T2
	ATOM	1978	CA	CYS	A	104	-5.378	61.978	-28.389	1.00	34.15	T2
	ATOM	1979	CB	CYS	A	104	-4.923	61.594	-26.987	1.00	29.38	T2
	ATOM	1980	SG	CYS	A	104	-5.706	62.501	-25.686	1.00	33.34	T2
	ATOM	1981	C	CYS	A	104	-4.888	63.367	-28.752	1.00	27.87	T2
20	ATOM	1982	O	CYS	A	104	-3.691	63.593	-28.921	1.00	34.61	T2
	ATOM	1983	N	TYR	A	105	-5.831	64.290	-28.888	1.00	27.29	T2
	ATOM	1984	CA	TYR	A	105	-5.529	65.670	-29.230	1.00	32.02	T2
	ATOM	1985	CB	TYR	A	105	-6.470	66.141	-30.335	1.00	31.54	T2
	ATOM	1986	CG	TYR	A	105	-6.331	67.600	-30.714	1.00	30.10	T2
25	ATOM	1987	CD1	TYR	A	105	-5.525	67.986	-31.777	1.00	30.24	T2
	ATOM	1988	CE1	TYR	A	105	-5.405	69.320	-32.138	1.00	31.35	T2
	ATOM	1989	CD2	TYR	A	105	-7.016	68.594	-30.013	1.00	27.39	T2
	ATOM	1990	CE2	TYR	A	105	-6.899	69.932	-30.361	1.00	27.60	T2
	ATOM	1991	CZ	TYR	A	105	-6.094	70.286	-31.427	1.00	24.99	T2
30	ATOM	1992	OH	TYR	A	105	-5.982	71.606	-31.800	1.00	22.31	T2
	ATOM	1993	C	TYR	A	105	-5.721	66.548	-28.008	1.00	27.98	T2
	ATOM	1994	O	TYR	A	105	-6.589	66.289	-27.181	1.00	33.26	T2
	ATOM	1995	N	SER	A	106	-4.909	67.588	-27.896	1.00	34.59	T2
	ATOM	1996	CA	SER	A	106	-5.021	68.516	-26.787	1.00	31.74	T2
35	ATOM	1997	CB	SER	A	106	-4.351	67.946	-25.531	1.00	27.43	T2
	ATOM	1998	OG	SER	A	106	-4.642	68.727	-24.386	1.00	32.45	T2
	ATOM	1999	C	SER	A	106	-4.340	69.797	-27.228	1.00	27.89	T2
	ATOM	2000	O	SER	A	106	-3.351	69.754	-27.959	1.00	31.90	T2
	ATOM	2001	N	ALA	A	107	-4.880	70.933	-26.797	1.00	24.08	T2
40	ATOM	2002	CA	ALA	A	107	-4.330	72.230	-27.163	1.00	38.96	T2
	ATOM	2003	CB	ALA	A	107	-4.840	72.637	-28.540	1.00	27.21	T2
	ATOM	2004	C	ALA	A	107	-4.694	73.294	-26.140	1.00	29.52	T2
	ATOM	2005	O	ALA	A	107	-5.583	73.093	-25.315	1.00	40.39	T2
	ATOM	2006	N	GLY	A	108	-4.001	74.423	-26.201	1.00	34.62	T2
45	ATOM	2007	CA	GLY	A	108	-4.268	75.506	-25.276	1.00	29.15	T2
	ATOM	2008	C	GLY	A	108	-3.494	76.751	-25.656	1.00	25.37	T2
	ATOM	2009	O	GLY	A	108	-2.756	76.744	-26.641	1.00	36.31	T2
	ATOM	2010	N	ILE	A	109	-3.661	77.822	-24.885	1.00	24.77	T2
	ATOM	2011	CA	ILE	A	109	-2.972	79.077	-25.150	1.00	28.20	T2
50	ATOM	2012	CB	ILE	A	109	-3.956	80.257	-25.200	1.00	31.51	T2
	ATOM	2013	CG2	ILE	A	109	-3.192	81.550	-25.420	1.00	28.63	T2
	ATOM	2014	CG1	ILE	A	109	-4.987	80.032	-26.308	1.00	36.22	T2
	ATOM	2015	CD1	ILE	A	109	-6.099	81.028	-26.309	1.00	21.38	T2
	ATOM	2016	C	ILE	A	109	-1.981	79.337	-24.035	1.00	25.51	T2
55	ATOM	2017	O	ILE	A	109	-2.265	79.064	-22.868	1.00	32.15	T2
	ATOM	2018	N	ALA	A	110	-0.819	79.865	-24.392	1.00	38.54	T2
	ATOM	2019	CA	ALA	A	110	0.209	80.161	-23.405	1.00	28.55	T2
	ATOM	2020	CB	ALA	A	110	1.114	78.963	-23.225	1.00	32.01	T2
	ATOM	2021	C	ALA	A	110	1.019	81.355	-23.868	1.00	30.86	T2
60	ATOM	2022	O	ALA	A	110	1.083	81.637	-25.064	1.00	27.53	T2
	ATOM	2023	N	LYS	A	111	1.624	82.073	-22.930	1.00	23.02	T2
	ATOM	2024	CA	LYS	A	111	2.449	83.205	-23.308	1.00	30.53	T2
	ATOM	2025	CB	LYS	A	111	2.337	84.326	-22.293	1.00	22.17	T2
	ATOM	2026	CG	LYS	A	111	3.152	85.547	-22.705	1.00	33.16	T2
65	ATOM	2027	CD	LYS	A	111	3.010	86.696	-21.709	1.00	23.93	T2
	ATOM	2028	CE	LYS	A	111	3.795	87.931	-22.168	1.00	35.50	T2

	ATOM	2029	NZ	LYS	A	111	3.627	89.086	-21.218	1.00	32.62	T2
	ATOM	2030	C	LYS	A	111	3.884	82.716	-23.376	1.00	33.39	T2
	ATOM	2031	O	LYS	A	111	4.349	82.039	-22.466	1.00	29.03	T2
	ATOM	2032	N	LEU	A	112	4.575	83.052	-24.460	1.00	33.89	T2
5	ATOM	2033	CA	LEU	A	112	5.957	82.625	-24.666	1.00	34.39	T2
	ATOM	2034	CB	LEU	A	112	6.021	81.627	-25.822	1.00	30.32	T2
	ATOM	2035	CG	LEU	A	112	5.029	80.470	-25.789	1.00	29.74	T2
	ATOM	2036	CD1	LEU	A	112	5.016	79.775	-27.136	1.00	27.93	T2
	ATOM	2037	CD2	LEU	A	112	5.394	79.512	-24.683	1.00	24.19	T2
10	ATOM	2038	C	LEU	A	112	6.854	83.813	-24.996	1.00	26.90	T2
	ATOM	2039	O	LEU	A	112	6.363	84.874	-25.390	1.00	29.44	T2
	ATOM	2040	N	GLU	A	113	8.167	83.628	-24.848	1.00	27.46	T2
	ATOM	2041	CA	GLU	A	113	9.133	84.685	-25.142	1.00	27.13	T2
	ATOM	2042	CB	GLU	A	113	9.825	85.175	-23.882	1.00	28.21	T2
15	ATOM	2043	CG	GLU	A	113	8.975	85.285	-22.656	1.00	30.48	T2
	ATOM	2044	CD	GLU	A	113	9.620	86.206	-21.625	1.00	31.84	T2
	ATOM	2045	OE1	GLU	A	113	10.860	86.070	-21.384	1.00	34.99	T2
	ATOM	2046	OE2	GLU	A	113	8.879	87.070	-21.060	1.00	28.50	T2
	ATOM	2047	C	GLU	A	113	10.231	84.230	-26.075	1.00	33.36	T2
20	ATOM	2048	O	GLU	A	113	10.520	83.044	-26.173	1.00	33.77	T2
	ATOM	2049	N	GLU	A	114	10.859	85.197	-26.735	1.00	30.17	T2
	ATOM	2050	CA	GLU	A	114	11.971	84.921	-27.637	1.00	32.43	T2
	ATOM	2051	CB	GLU	A	114	12.787	86.175	-27.894	1.00	25.29	T2
	ATOM	2052	CG	GLU	A	114	12.357	87.039	-29.043	1.00	29.47	T2
25	ATOM	2053	CD	GLU	A	114	13.491	87.963	-29.464	1.00	29.06	T2
	ATOM	2054	OE1	GLU	A	114	14.549	87.439	-29.898	1.00	33.95	T2
	ATOM	2055	OE2	GLU	A	114	13.329	89.199	-29.346	1.00	30.05	T2
	ATOM	2056	C	GLU	A	114	12.924	83.942	-26.984	1.00	30.90	T2
	ATOM	2057	O	GLU	A	114	13.417	84.197	-25.885	1.00	31.10	T2
30	ATOM	2058	N	GLY	A	115	13.214	82.841	-27.659	1.00	32.95	T2
	ATOM	2059	CA	GLY	A	115	14.141	81.891	-27.089	1.00	36.50	T2
	ATOM	2060	C	GLY	A	115	13.490	80.701	-26.426	1.00	33.62	T2
	ATOM	2061	O	GLY	A	115	14.152	79.694	-26.191	1.00	36.09	T2
	ATOM	2062	N	ASP	A	116	12.211	80.806	-26.086	1.00	35.38	T2
35	ATOM	2063	CA	ASP	A	116	11.548	79.668	-25.473	1.00	24.69	T2
	ATOM	2064	CB	ASP	A	116	10.128	80.022	-25.020	1.00	37.12	T2
	ATOM	2065	CG	ASP	A	116	10.100	80.941	-23.816	1.00	26.84	T2
	ATOM	2066	OD1	ASP	A	116	11.077	80.931	-23.042	1.00	28.57	T2
	ATOM	2067	OD2	ASP	A	116	9.092	81.661	-23.628	1.00	31.42	T2
40	ATOM	2068	C	ASP	A	116	11.472	78.564	-26.525	1.00	36.51	T2
	ATOM	2069	O	ASP	A	116	11.466	78.834	-27.730	1.00	31.46	T2
	ATOM	2070	N	GLU	A	117	11.431	77.319	-26.072	1.00	33.81	T2
	ATOM	2071	CA	GLU	A	117	11.328	76.203	-26.994	1.00	33.70	T2
	ATOM	2072	CB	GLU	A	117	12.607	75.368	-26.988	1.00	33.57	T2
45	ATOM	2073	CG	GLU	A	117	13.860	76.160	-27.267	1.00	27.95	T2
	ATOM	2074	CD	GLU	A	117	15.095	75.282	-27.338	1.00	30.73	T2
	ATOM	2075	OE1	GLU	A	117	15.171	74.307	-26.554	1.00	31.98	T2
	ATOM	2076	OE2	GLU	A	117	15.990	75.576	-28.165	1.00	30.83	T2
	ATOM	2077	C	GLU	A	117	10.160	75.335	-26.568	1.00	29.80	T2
50	ATOM	2078	O	GLU	A	117	9.929	75.135	-25.369	1.00	29.86	T2
	ATOM	2079	N	LEU	A	118	9.416	74.835	-27.552	1.00	25.19	T2
	ATOM	2080	CA	LEU	A	118	8.281	73.966	-27.285	1.00	31.74	T2
	ATOM	2081	CB	LEU	A	118	7.048	74.426	-28.052	1.00	34.81	T2
	ATOM	2082	CG	LEU	A	118	6.462	75.776	-27.663	1.00	28.70	T2
55	ATOM	2083	CD1	LEU	A	118	5.209	76.026	-28.465	1.00	32.69	T2
	ATOM	2084	CD2	LEU	A	118	6.149	75.789	-26.186	1.00	28.34	T2
	ATOM	2085	C	LEU	A	118	8.648	72.574	-27.740	1.00	33.22	T2
	ATOM	2086	O	LEU	A	118	9.367	72.416	-28.724	1.00	29.66	T2
	ATOM	2087	N	GLN	A	119	8.159	71.568	-27.020	1.00	30.73	T2
60	ATOM	2088	CA	GLN	A	119	8.428	70.184	-27.379	1.00	32.03	T2
	ATOM	2089	CB	GLN	A	119	9.741	69.730	-26.758	1.00	29.73	T2
	ATOM	2090	CG	GLN	A	119	9.737	69.738	-25.256	1.00	26.55	T2
	ATOM	2091	CD	GLN	A	119	11.086	69.359	-24.675	1.00	25.11	T2
	ATOM	2092	OE1	GLN	A	119	11.192	69.001	-23.497	1.00	29.14	T2
65	ATOM	2093	NE2	GLN	A	119	12.131	69.446	-25.494	1.00	31.14	T2
	ATOM	2094	C	GLN	A	119	7.296	69.264	-26.944	1.00	29.35	T2

	ATOM	2095	O	GLN	A	119	6.566	69.570	-26.005	1.00	29.09	T2
	ATOM	2096	N	LEU	A	120	7.161	68.141	-27.641	1.00	28.13	T2
	ATOM	2097	CA	LEU	A	120	6.123	67.146	-27.376	1.00	33.73	T2
	ATOM	2098	CB	LEU	A	120	5.428	66.789	-28.698	1.00	29.77	T2
5	ATOM	2099	CG	LEU	A	120	4.127	65.991	-28.826	1.00	31.02	T2
	ATOM	2100	CD1	LEU	A	120	4.154	64.766	-27.927	1.00	24.71	T2
	ATOM	2101	CD2	LEU	A	120	2.971	66.884	-28.485	1.00	29.37	T2
	ATOM	2102	C	LEU	A	120	6.823	65.918	-26.806	1.00	38.44	T2
	ATOM	2103	O	LEU	A	120	7.688	65.342	-27.463	1.00	25.46	T2
10	ATOM	2104	N	ALA	A	121	6.457	65.510	-25.596	1.00	25.95	T2
	ATOM	2105	CA	ALA	A	121	7.105	64.353	-24.987	1.00	30.95	T2
	ATOM	2106	CB	ALA	A	121	8.009	64.822	-23.868	1.00	35.27	T2
	ATOM	2107	C	ALA	A	121	6.171	63.257	-24.468	1.00	22.80	T2
	ATOM	2108	O	ALA	A	121	5.108	63.534	-23.902	1.00	22.16	T2
15	ATOM	2109	N	ILE	A	122	6.585	62.007	-24.666	1.00	33.79	T2
	ATOM	2110	CA	ILE	A	122	5.824	60.855	-24.191	1.00	31.14	T2
	ATOM	2111	CB	ILE	A	122	5.672	59.788	-25.275	1.00	33.44	T2
	ATOM	2112	CG2	ILE	A	122	4.836	58.641	-24.740	1.00	24.78	T2
	ATOM	2113	CG1	ILE	A	122	4.998	60.397	-26.504	1.00	33.19	T2
20	ATOM	2114	CD1	ILE	A	122	4.795	59.434	-27.647	1.00	34.27	T2
	ATOM	2115	C	ILE	A	122	6.572	60.245	-23.008	1.00	23.27	T2
	ATOM	2116	O	ILE	A	122	7.712	59.786	-23.145	1.00	28.80	T2
	ATOM	2117	N	PRO	A	123	5.941	60.241	-21.826	1.00	30.24	T2
	ATOM	2118	CD	PRO	A	123	4.644	60.879	-21.569	1.00	28.92	T2
25	ATOM	2119	CA	PRO	A	123	6.509	59.704	-20.581	1.00	36.16	T2
	ATOM	2120	CB	PRO	A	123	5.555	60.226	-19.506	1.00	35.36	T2
	ATOM	2121	CG	PRO	A	123	4.846	61.385	-20.173	1.00	31.58	T2
	ATOM	2122	C	PRO	A	123	6.605	58.179	-20.542	1.00	34.03	T2
	ATOM	2123	O	PRO	A	123	6.072	57.536	-19.631	1.00	31.27	T2
30	ATOM	2124	N	ARG	A	124	7.271	57.602	-21.536	1.00	30.01	T2
	ATOM	2125	CA	ARG	A	124	7.453	56.156	-21.604	1.00	20.56	T2
	ATOM	2126	CB	ARG	A	124	6.334	55.491	-22.380	1.00	32.63	T2
	ATOM	2127	CG	ARG	A	124	5.136	55.183	-21.529	1.00	29.78	T2
	ATOM	2128	CD	ARG	A	124	4.642	53.794	-21.881	1.00	32.83	T2
35	ATOM	2129	NE	ARG	A	124	5.532	52.744	-21.374	1.00	35.07	T2
	ATOM	2130	CZ	ARG	A	124	5.409	51.457	-21.690	1.00	29.64	T2
	ATOM	2131	NH1	ARG	A	124	4.440	51.077	-22.521	1.00	26.52	T2
	ATOM	2132	NH2	ARG	A	124	6.231	50.551	-21.165	1.00	24.67	T2
	ATOM	2133	C	ARG	A	124	8.760	55.862	-22.282	1.00	33.82	T2
40	ATOM	2134	O	ARG	A	124	9.261	56.679	-23.056	1.00	29.88	T2
	ATOM	2135	N	GLU	A	125	9.307	54.685	-22.008	1.00	31.73	T2
	ATOM	2136	CA	GLU	A	125	10.595	54.333	-22.585	1.00	37.55	T2
	ATOM	2137	CB	GLU	A	125	11.224	53.178	-21.811	1.00	30.87	T2
	ATOM	2138	CG	GLU	A	125	11.715	53.612	-20.430	1.00	41.01	T2
45	ATOM	2139	CD	GLU	A	125	13.069	53.004	-20.080	1.00	25.88	T2
	ATOM	2140	OE1	GLU	A	125	13.178	51.743	-20.095	1.00	31.14	T2
	ATOM	2141	OE2	GLU	A	125	14.020	53.789	-19.793	1.00	29.48	T2
	ATOM	2142	C	GLU	A	125	10.573	54.036	-24.072	1.00	29.94	T2
	ATOM	2143	O	GLU	A	125	11.386	54.584	-24.817	1.00	28.97	T2
50	ATOM	2144	N	ASN	A	126	9.676	53.169	-24.521	1.00	25.45	T2
	ATOM	2145	CA	ASN	A	126	9.625	52.904	-25.952	1.00	26.95	T2
	ATOM	2146	CB	ASN	A	126	10.329	51.592	-26.311	1.00	28.03	T2
	ATOM	2147	CG	ASN	A	126	11.846	51.757	-26.415	1.00	34.00	T2
	ATOM	2148	OD1	ASN	A	126	12.554	51.806	-25.400	1.00	25.98	T2
55	ATOM	2149	ND2	ASN	A	126	12.347	51.864	-27.646	1.00	30.41	T2
	ATOM	2150	C	ASN	A	126	8.188	52.866	-26.382	1.00	28.80	T2
	ATOM	2151	O	ASN	A	126	7.680	51.839	-26.846	1.00	28.61	T2
	ATOM	2152	N	ALA	A	127	7.537	54.012	-26.214	1.00	35.04	T2
	ATOM	2153	CA	ALA	A	127	6.135	54.160	-26.553	1.00	30.90	T2
60	ATOM	2154	CB	ALA	A	127	5.755	55.626	-26.499	1.00	30.02	T2
	ATOM	2155	C	ALA	A	127	5.810	53.584	-27.921	1.00	29.58	T2
	ATOM	2156	O	ALA	A	127	6.516	53.846	-28.888	1.00	31.25	T2
	ATOM	2157	N	GLN	A	128	4.755	52.776	-27.980	1.00	29.02	T2
	ATOM	2158	CA	GLN	A	128	4.307	52.200	-29.241	1.00	27.77	T2
65	ATOM	2159	CB	GLN	A	128	3.536	50.904	-28.991	1.00	25.47	T2
	ATOM	2160	CG	GLN	A	128	4.424	49.795	-28.433	1.00	33.16	T2

	ATOM	2161	CD	GLN	A	128	5.735	49.678	-29.202	1.00	37.27	T2
	ATOM	2162	OE1	GLN	A	128	5.742	49.452	-30.416	1.00	33.24	T2
	ATOM	2163	NE2	GLN	A	128	6.850	49.842	-28.499	1.00	32.61	T2
	ATOM	2164	C	GLN	A	128	3.409	53.262	-29.877	1.00	27.55	T2
5	ATOM	2165	O	GLN	A	128	2.234	53.413	-29.526	1.00	31.55	T2
	ATOM	2166	N	ILE	A	129	3.987	53.989	-30.827	1.00	30.63	T2
	ATOM	2167	CA	ILE	A	129	3.331	55.104	-31.481	1.00	32.51	T2
	ATOM	2168	CB	ILE	A	129	4.140	56.402	-31.114	1.00	27.33	T2
	ATOM	2169	CG2	ILE	A	129	4.505	57.210	-32.333	1.00	32.11	T2
10	ATOM	2170	CG1	ILE	A	129	3.368	57.204	-30.079	1.00	29.84	T2
	ATOM	2171	CD1	ILE	A	129	3.127	56.433	-28.803	1.00	24.08	T2
	ATOM	2172	C	ILE	A	129	3.215	54.947	-32.988	1.00	34.73	T2
	ATOM	2173	O	ILE	A	129	3.946	54.171	-33.591	1.00	28.89	T2
	ATOM	2174	N	SER	A	130	2.280	55.672	-33.590	1.00	33.79	T2
15	ATOM	2175	CA	SER	A	130	2.128	55.644	-35.042	1.00	29.28	T2
	ATOM	2176	CB	SER	A	130	0.652	55.640	-35.442	1.00	29.94	T2
	ATOM	2177	OG	SER	A	130	0.512	55.885	-36.839	1.00	27.55	T2
	ATOM	2178	C	SER	A	130	2.786	56.912	-35.581	1.00	29.78	T2
	ATOM	2179	O	SER	A	130	2.459	58.012	-35.136	1.00	33.26	T2
20	ATOM	2180	N	LEU	A	131	3.708	56.777	-36.526	1.00	32.06	T2
	ATOM	2181	CA	LEU	A	131	4.366	57.959	-37.066	1.00	32.91	T2
	ATOM	2182	CB	LEU	A	131	5.861	57.710	-37.245	1.00	27.46	T2
	ATOM	2183	CG	LEU	A	131	6.699	57.802	-35.967	1.00	25.92	T2
	ATOM	2184	CD1	LEU	A	131	6.217	56.799	-34.950	1.00	28.55	T2
25	ATOM	2185	CD2	LEU	A	131	8.151	57.535	-36.302	1.00	24.65	T2
	ATOM	2186	C	LEU	A	131	3.781	58.467	-38.377	1.00	28.20	T2
	ATOM	2187	O	LEU	A	131	4.510	58.993	-39.223	1.00	29.19	T2
	ATOM	2188	N	ASP	A	132	2.470	58.323	-38.548	1.00	26.30	T2
	ATOM	2189	CA	ASP	A	132	1.817	58.799	-39.761	1.00	27.83	T2
30	ATOM	2190	CB	ASP	A	132	0.570	57.965	-40.059	1.00	29.45	T2
	ATOM	2191	CG	ASP	A	132	0.893	56.666	-40.790	1.00	31.15	T2
	ATOM	2192	OD1	ASP	A	132	-0.004	55.795	-40.889	1.00	30.95	T2
	ATOM	2193	OD2	ASP	A	132	2.041	56.523	-41.274	1.00	30.04	T2
	ATOM	2194	C	ASP	A	132	1.443	60.279	-39.678	1.00	32.84	T2
35	ATOM	2195	O	ASP	A	132	1.013	60.775	-38.630	1.00	27.33	T2
	ATOM	2196	N	GLY	A	133	1.612	60.972	-40.802	1.00	26.48	T2
	ATOM	2197	CA	GLY	A	133	1.304	62.386	-40.871	1.00	31.46	T2
	ATOM	2198	C	GLY	A	133	-0.089	62.768	-40.411	1.00	22.43	T2
	ATOM	2199	O	GLY	A	133	-0.258	63.857	-39.878	1.00	26.76	T2
40	ATOM	2200	N	ASP	A	134	-1.084	61.899	-40.606	1.00	33.94	T2
	ATOM	2201	CA	ASP	A	134	-2.449	62.223	-40.193	1.00	25.07	T2
	ATOM	2202	CB	ASP	A	134	-3.471	61.200	-40.664	1.00	35.87	T2
	ATOM	2203	CG	ASP	A	134	-3.048	60.473	-41.865	1.00	27.08	T2
	ATOM	2204	OD1	ASP	A	134	-2.897	61.140	-42.900	1.00	27.02	T2
45	ATOM	2205	OD2	ASP	A	134	-2.877	59.238	-41.771	1.00	29.61	T2
	ATOM	2206	C	ASP	A	134	-2.590	62.204	-38.701	1.00	28.12	T2
	ATOM	2207	O	ASP	A	134	-2.998	63.176	-38.081	1.00	37.65	T2
	ATOM	2208	N	VAL	A	135	-2.278	61.048	-38.138	1.00	28.54	T2
	ATOM	2209	CA	VAL	A	135	-2.436	60.814	-36.719	1.00	33.88	T2
50	ATOM	2210	CB	VAL	A	135	-2.351	59.322	-36.459	1.00	31.50	T2
	ATOM	2211	CG1	VAL	A	135	-3.464	58.629	-37.231	1.00	34.89	T2
	ATOM	2212	CG2	VAL	A	135	-0.996	58.796	-36.898	1.00	34.65	T2
	ATOM	2213	C	VAL	A	135	-1.589	61.561	-35.695	1.00	22.76	T2
	ATOM	2214	O	VAL	A	135	-2.109	61.915	-34.636	1.00	34.49	T2
55	ATOM	2215	N	THR	A	136	-0.309	61.808	-35.963	1.00	30.25	T2
	ATOM	2216	CA	THR	A	136	0.475	62.534	-34.965	1.00	32.47	T2
	ATOM	2217	CB	THR	A	136	1.493	61.597	-34.254	1.00	33.95	T2
	ATOM	2218	OG1	THR	A	136	2.619	61.367	-35.098	1.00	30.99	T2
	ATOM	2219	CG2	THR	A	136	0.853	60.259	-33.946	1.00	31.56	T2
60	ATOM	2220	C	THR	A	136	1.176	63.785	-35.515	1.00	26.98	T2
	ATOM	2221	O	THR	A	136	1.982	63.716	-36.443	1.00	33.61	T2
	ATOM	2222	N	PHE	A	137	0.841	64.934	-34.930	1.00	24.10	T2
	ATOM	2223	CA	PHE	A	137	1.392	66.216	-35.351	1.00	30.39	T2
	ATOM	2224	CB	PHE	A	137	0.485	66.819	-36.419	1.00	24.19	T2
65	ATOM	2225	CG	PHE	A	137	-0.983	66.718	-36.103	1.00	26.74	T2
	ATOM	2226	CD1	PHE	A	137	-1.587	67.615	-35.241	1.00	32.52	T2

	ATOM	2227	CD2	PHE	A	137	-1.759	65.724	-36.675	1.00	39.60	T2
	ATOM	2228	CE1	PHE	A	137	-2.940	67.525	-34.957	1.00	28.07	T2
	ATOM	2229	CE2	PHE	A	137	-3.108	65.628	-36.395	1.00	39.80	T2
	ATOM	2230	CZ	PHE	A	137	-3.698	66.529	-35.536	1.00	31.01	T2
5	ATOM	2231	C	PHE	A	137	1.574	67.189	-34.185	1.00	32.98	T2
	ATOM	2232	O	PHE	A	137	1.017	66.992	-33.110	1.00	33.16	T2
	ATOM	2233	N	PHE	A	138	2.350	68.245	-34.412	1.00	29.40	T2
	ATOM	2234	CA	PHE	A	138	2.641	69.219	-33.366	1.00	27.26	T2
	ATOM	2235	CB	PHE	A	138	4.106	69.032	-32.947	1.00	30.44	T2
10	ATOM	2236	CG	PHE	A	138	4.478	69.718	-31.669	1.00	27.93	T2
	ATOM	2237	CD1	PHE	A	138	3.515	70.032	-30.720	1.00	31.33	T2
	ATOM	2238	CD2	PHE	A	138	5.805	70.046	-31.415	1.00	39.23	T2
	ATOM	2239	CE1	PHE	A	138	3.868	70.669	-29.535	1.00	30.38	T2
	ATOM	2240	CE2	PHE	A	138	6.169	70.679	-30.238	1.00	29.74	T2
15	ATOM	2241	CZ	PHE	A	138	5.201	70.993	-29.295	1.00	29.01	T2
	ATOM	2242	C	PHE	A	138	2.332	70.668	-33.799	1.00	29.04	T2
	ATOM	2243	O	PHE	A	138	2.756	71.127	-34.860	1.00	32.39	T2
	ATOM	2244	N	GLY	A	139	1.602	71.369	-32.928	1.00	32.54	T2
	ATOM	2245	CA	GLY	A	139	1.115	72.724	-33.162	1.00	28.06	T2
20	ATOM	2246	C	GLY	A	139	1.913	74.008	-33.175	1.00	32.38	T2
	ATOM	2247	O	GLY	A	139	2.948	74.066	-33.808	1.00	36.20	T2
	ATOM	2248	N	ALA	A	140	1.376	75.044	-32.520	1.00	27.54	T2
	ATOM	2249	CA	ALA	A	140	1.963	76.394	-32.412	1.00	28.16	T2
	ATOM	2250	CB	ALA	A	140	3.459	76.310	-32.343	1.00	32.60	T2
25	ATOM	2251	C	ALA	A	140	1.569	77.418	-33.482	1.00	30.02	T2
	ATOM	2252	O	ALA	A	140	2.043	77.370	-34.609	1.00	31.34	T2
	ATOM	2253	N	LEU	A	141	0.714	78.358	-33.093	1.00	35.48	T2
	ATOM	2254	CA	LEU	A	141	0.228	79.439	-33.960	1.00	33.58	T2
	ATOM	2255	CB	LEU	A	141	-1.174	79.111	-34.487	1.00	29.23	T2
30	ATOM	2256	CG	LEU	A	141	-1.955	80.175	-35.266	1.00	32.24	T2
	ATOM	2257	CD1	LEU	A	141	-3.069	79.530	-36.030	1.00	25.82	T2
	ATOM	2258	CD2	LEU	A	141	-2.518	81.194	-34.325	1.00	30.57	T2
	ATOM	2259	C	LEU	A	141	0.175	80.724	-33.137	1.00	30.21	T2
	ATOM	2260	O	LEU	A	141	-0.317	80.720	-32.009	1.00	34.22	T2
35	ATOM	2261	N	LYS	A	142	0.660	81.826	-33.695	1.00	32.81	T2
	ATOM	2262	CA	LYS	A	142	0.668	83.084	-32.953	1.00	28.98	T2
	ATOM	2263	CB	LYS	A	142	1.867	83.936	-33.377	1.00	29.29	T2
	ATOM	2264	CG	LYS	A	142	1.938	85.245	-32.628	1.00	30.34	T2
	ATOM	2265	CD	LYS	A	142	3.255	85.959	-32.810	1.00	32.15	T2
40	ATOM	2266	CE	LYS	A	142	3.253	87.249	-32.004	1.00	26.24	T2
	ATOM	2267	NZ	LYS	A	142	4.509	88.013	-32.183	1.00	27.83	T2
	ATOM	2268	C	LYS	A	142	-0.607	83.932	-33.041	1.00	24.56	T2
	ATOM	2269	O	LYS	A	142	-1.087	84.242	-34.131	1.00	30.25	T2
	ATOM	2270	N	LEU	A	143	-1.142	84.319	-31.887	1.00	33.17	T2
45	ATOM	2271	CA	LEU	A	143	-2.346	85.136	-31.836	1.00	30.78	T2
	ATOM	2272	CB	LEU	A	143	-3.021	84.994	-30.477	1.00	33.44	T2
	ATOM	2273	CG	LEU	A	143	-3.440	83.599	-30.015	1.00	28.20	T2
	ATOM	2274	CD1	LEU	A	143	-4.008	83.689	-28.606	1.00	30.48	T2
	ATOM	2275	CD2	LEU	A	143	-4.472	83.024	-30.960	1.00	36.58	T2
50	ATOM	2276	C	LEU	A	143	-2.001	86.602	-32.061	1.00	24.11	T2
	ATOM	2277	O	LEU	A	143	-0.877	87.019	-31.807	1.00	26.20	T2
	ATOM	2278	N	LEU	A	144	-2.968	87.386	-32.532	1.00	23.08	T2
	ATOM	2279	CA	LEU	A	144	-2.738	88.807	-32.766	1.00	31.74	T2
	ATOM	2280	CB	LEU	A	144	-3.642	89.319	-33.881	1.00	33.13	T2
55	ATOM	2281	CG	LEU	A	144	-3.367	88.747	-35.275	1.00	26.98	T2
	ATOM	2282	CD1	LEU	A	144	-4.454	89.224	-36.230	1.00	24.72	T2
	ATOM	2283	CD2	LEU	A	144	-1.985	89.183	-35.773	1.00	26.04	T2
	ATOM	2284	C	LEU	A	144	-3.015	89.594	-31.500	1.00	29.20	T2
	ATOM	2285	O	LEU	A	144	-3.686	89.041	-30.603	1.00	32.70	T2
60	ATOM	2286	OXT	LEU	A	144	-2.569	90.759	-31.424	1.00	28.62	T2
	ATOM	2287	CB	VAL	A	1	-5.368	96.201	-32.612	1.00	32.33	T3
	ATOM	2288	CG1	VAL	A	1	-3.821	96.164	-32.639	1.00	34.91	T3
	ATOM	2289	CG2	VAL	A	1	-5.875	97.537	-33.169	1.00	29.59	T3
	ATOM	2290	C	VAL	A	1	-5.518	93.677	-32.799	1.00	31.75	T3
65	ATOM	2291	O	VAL	A	1	-4.813	92.885	-33.435	1.00	25.86	T3
	ATOM	2292	N	VAL	A	1	-7.446	95.103	-33.573	1.00	32.89	T3

	ATOM	2293	CA	VAL	A	1	-5.958	95.006	-33.449	1.00	28.48	T3
	ATOM	2294	N	THR	A	2	-5.925	93.432	-31.549	1.00	29.53	T3
	ATOM	2295	CA	THR	A	2	-5.537	92.190	-30.871	1.00	26.61	T3
	ATOM	2296	CB	THR	A	2	-4.859	92.464	-29.516	1.00	35.08	T3
5	ATOM	2297	OG1	THR	A	2	-5.842	92.936	-28.581	1.00	31.47	T3
	ATOM	2298	CG2	THR	A	2	-3.748	93.497	-29.672	1.00	34.97	T3
	ATOM	2299	C	THR	A	2	-6.715	91.255	-30.609	1.00	30.93	T3
	ATOM	2300	O	THR	A	2	-7.868	91.627	-30.807	1.00	32.86	T3
	ATOM	2301	N	GLN	A	3	-6.414	90.043	-30.147	1.00	25.63	T3
10	ATOM	2302	CA	GLN	A	3	-7.442	89.050	-29.851	1.00	27.13	T3
	ATOM	2303	CB	GLN	A	3	-7.138	87.742	-30.577	1.00	38.73	T3
	ATOM	2304	CG	GLN	A	3	-6.757	87.908	-32.026	1.00	32.08	T3
	ATOM	2305	CD	GLN	A	3	-6.616	86.575	-32.721	1.00	24.82	T3
	ATOM	2306	OE1	GLN	A	3	-7.567	85.803	-32.789	1.00	29.27	T3
15	ATOM	2307	NE2	GLN	A	3	-5.426	86.295	-33.237	1.00	36.82	T3
	ATOM	2308	C	GLN	A	3	-7.545	88.758	-28.360	1.00	33.67	T3
	ATOM	2309	O	GLN	A	3	-6.707	88.049	-27.805	1.00	29.02	T3
	ATOM	2310	N	ASP	A	4	-8.574	89.287	-27.712	1.00	34.10	T3
	ATOM	2311	CA	ASP	A	4	-8.735	89.037	-26.295	1.00	27.52	T3
20	ATOM	2312	CB	ASP	A	4	-9.944	89.794	-25.751	1.00	33.10	T3
	ATOM	2313	CG	ASP	A	4	-9.784	91.294	-25.855	1.00	34.28	T3
	ATOM	2314	OD1	ASP	A	4	-8.635	91.746	-26.028	1.00	33.67	T3
	ATOM	2315	OD2	ASP	A	4	-10.797	92.023	-25.754	1.00	29.48	T3
	ATOM	2316	C	ASP	A	4	-8.910	87.547	-26.046	1.00	30.37	T3
25	ATOM	2317	O	ASP	A	4	-9.466	86.842	-26.882	1.00	29.28	T3
	ATOM	2318	N	CYS	A	5	-8.421	87.074	-24.901	1.00	29.91	T3
	ATOM	2319	CA	CYS	A	5	-8.535	85.669	-24.519	1.00	37.37	T3
	ATOM	2320	CB	CYS	A	5	-7.456	84.813	-25.208	1.00	30.06	T3
	ATOM	2321	SG	CYS	A	5	-5.871	85.632	-25.596	1.00	34.41	T3
30	ATOM	2322	C	CYS	A	5	-8.423	85.533	-23.009	1.00	26.17	T3
	ATOM	2323	O	CYS	A	5	-7.736	86.312	-22.361	1.00	30.61	T3
	ATOM	2324	N	LEU	A	6	-9.128	84.562	-22.444	1.00	38.75	T3
	ATOM	2325	CA	LEU	A	6	-9.087	84.323	-21.005	1.00	31.71	T3
	ATOM	2326	CB	LEU	A	6	-10.323	84.905	-20.321	1.00	30.51	T3
35	ATOM	2327	CG	LEU	A	6	-10.444	84.685	-18.812	1.00	33.61	T3
	ATOM	2328	CD1	LEU	A	6	-11.206	85.834	-18.179	1.00	26.46	T3
	ATOM	2329	CD2	LEU	A	6	-11.146	83.375	-18.550	1.00	29.34	T3
	ATOM	2330	C	LEU	A	6	-9.041	82.827	-20.808	1.00	34.84	T3
	ATOM	2331	O	LEU	A	6	-9.794	82.099	-21.439	1.00	27.75	T3
40	ATOM	2332	N	GLN	A	7	-8.156	82.363	-19.935	1.00	36.79	T3
	ATOM	2333	CA	GLN	A	7	-8.023	80.934	-19.703	1.00	38.40	T3
	ATOM	2334	CB	GLN	A	7	-6.718	80.438	-20.314	1.00	33.10	T3
	ATOM	2335	CG	GLN	A	7	-6.581	78.941	-20.368	1.00	29.78	T3
	ATOM	2336	CD	GLN	A	7	-5.384	78.528	-21.191	1.00	28.27	T3
45	ATOM	2337	OE1	GLN	A	7	-4.240	78.741	-20.797	1.00	32.06	T3
	ATOM	2338	NE2	GLN	A	7	-5.640	77.953	-22.353	1.00	29.18	T3
	ATOM	2339	C	GLN	A	7	-8.063	80.590	-18.229	1.00	25.22	T3
	ATOM	2340	O	GLN	A	7	-7.517	81.309	-17.398	1.00	30.36	T3
	ATOM	2341	N	LEU	A	8	-8.718	79.480	-17.917	1.00	31.59	T3
50	ATOM	2342	CA	LEU	A	8	-8.843	79.020	-16.544	1.00	30.76	T3
	ATOM	2343	CB	LEU	A	8	-10.323	78.894	-16.168	1.00	30.20	T3
	ATOM	2344	CG	LEU	A	8	-11.098	80.100	-15.636	1.00	28.42	T3
	ATOM	2345	CD1	LEU	A	8	-10.345	81.387	-15.881	1.00	31.53	T3
	ATOM	2346	CD2	LEU	A	8	-12.453	80.124	-16.294	1.00	33.95	T3
55	ATOM	2347	C	LEU	A	8	-8.142	77.681	-16.330	1.00	34.25	T3
	ATOM	2348	O	LEU	A	8	-7.991	76.887	-17.260	1.00	20.38	T3
	ATOM	2349	N	ILE	A	9	-7.726	77.444	-15.091	1.00	34.94	T3
	ATOM	2350	CA	ILE	A	9	-7.031	76.224	-14.699	1.00	33.67	T3
	ATOM	2351	CB	ILE	A	9	-5.584	76.541	-14.320	1.00	30.64	T3
60	ATOM	2352	CG2	ILE	A	9	-4.999	75.424	-13.493	1.00	29.30	T3
	ATOM	2353	CG1	ILE	A	9	-4.738	76.739	-15.559	1.00	24.57	T3
	ATOM	2354	CD1	ILE	A	9	-3.314	77.080	-15.189	1.00	31.91	T3
	ATOM	2355	C	ILE	A	9	-7.699	75.619	-13.468	1.00	33.76	T3
	ATOM	2356	O	ILE	A	9	-8.143	76.341	-12.588	1.00	24.33	T3
65	ATOM	2357	N	ALA	A	10	-7.747	74.298	-13.391	1.00	25.01	T3
	ATOM	2358	CA	ALA	A	10	-8.354	73.667	-12.232	1.00	28.75	T3

	ATOM	2359	CB	ALA	A	10	-8.349	72.160	-12.395	1.00	28.12	T3
	ATOM	2360	C	ALA	A	10	-7.616	74.050	-10.946	1.00	28.72	T3
	ATOM	2361	O	ALA	A	10	-6.380	73.972	-10.869	1.00	31.31	T3
	ATOM	2362	N	ASP	A	11	-8.377	74.457	-9.932	1.00	34.17	T3
5	ATOM	2363	CA	ASP	A	11	-7.792	74.829	-8.650	1.00	31.18	T3
	ATOM	2364	CB	ASP	A	11	-8.593	75.951	-7.992	1.00	30.66	T3
	ATOM	2365	CG	ASP	A	11	-8.094	76.266	-6.598	1.00	30.33	T3
	ATOM	2366	OD1	ASP	A	11	-6.863	76.224	-6.398	1.00	28.63	T3
	ATOM	2367	OD2	ASP	A	11	-8.922	76.553	-5.706	1.00	31.62	T3
10	ATOM	2368	C	ASP	A	11	-7.752	73.613	-7.739	1.00	34.84	T3
	ATOM	2369	O	ASP	A	11	-8.701	73.333	-7.002	1.00	31.56	T3
	ATOM	2370	N	SER	A	12	-6.633	72.900	-7.812	1.00	32.01	T3
	ATOM	2371	CA	SER	A	12	-6.406	71.683	-7.042	1.00	27.54	T3
	ATOM	2372	CB	SER	A	12	-5.086	71.020	-7.477	1.00	34.04	T3
15	ATOM	2373	OG	SER	A	12	-3.974	71.902	-7.327	1.00	29.04	T3
	ATOM	2374	C	SER	A	12	-6.374	71.911	-5.538	1.00	28.55	T3
	ATOM	2375	O	SER	A	12	-5.804	71.102	-4.799	1.00	29.06	T3
	ATOM	2376	N	GLU	A	13	-6.979	73.000	-5.075	1.00	32.41	T3
	ATOM	2377	CA	GLU	A	13	-6.978	73.263	-3.645	1.00	30.23	T3
20	ATOM	2378	CB	GLU	A	13	-5.892	74.259	-3.294	1.00	34.31	T3
	ATOM	2379	CG	GLU	A	13	-4.594	73.566	-2.985	1.00	29.35	T3
	ATOM	2380	CD	GLU	A	13	-3.531	74.551	-2.558	1.00	32.76	T3
	ATOM	2381	OE1	GLU	A	13	-3.876	75.476	-1.773	1.00	33.74	T3
	ATOM	2382	OE2	GLU	A	13	-2.355	74.402	-2.998	1.00	29.68	T3
25	ATOM	2383	C	GLU	A	13	-8.287	73.696	-3.032	1.00	38.53	T3
	ATOM	2384	O	GLU	A	13	-8.308	74.358	-1.990	1.00	37.15	T3
	ATOM	2385	N	THR	A	14	-9.379	73.326	-3.691	1.00	25.77	T3
	ATOM	2386	CA	THR	A	14	-10.708	73.612	-3.185	1.00	33.43	T3
	ATOM	2387	CB	THR	A	14	-11.282	74.944	-3.710	1.00	28.60	T3
30	ATOM	2388	OG1	THR	A	14	-11.277	74.925	-5.132	1.00	32.90	T3
	ATOM	2389	CG2	THR	A	14	-10.457	76.127	-3.226	1.00	21.91	T3
	ATOM	2390	C	THR	A	14	-11.561	72.458	-3.667	1.00	32.63	T3
	ATOM	2391	O	THR	A	14	-11.276	71.843	-4.696	1.00	29.90	T3
	ATOM	2392	N	PRO	A	15	-12.606	72.126	-2.910	1.00	26.58	T3
35	ATOM	2393	CD	PRO	A	15	-13.033	72.800	-1.677	1.00	27.56	T3
	ATOM	2394	CA	PRO	A	15	-13.517	71.029	-3.251	1.00	32.32	T3
	ATOM	2395	CB	PRO	A	15	-14.503	71.021	-2.085	1.00	30.30	T3
	ATOM	2396	CG	PRO	A	15	-13.738	71.695	-0.967	1.00	25.14	T3
	ATOM	2397	C	PRO	A	15	-14.230	71.291	-4.570	1.00	29.69	T3
40	ATOM	2398	O	PRO	A	15	-14.549	72.436	-4.894	1.00	27.61	T3
	ATOM	2399	N	THR	A	16	-14.486	70.237	-5.325	1.00	26.78	T3
	ATOM	2400	CA	THR	A	16	-15.189	70.396	-6.591	1.00	30.89	T3
	ATOM	2401	CB	THR	A	16	-15.139	69.096	-7.392	1.00	30.83	T3
	ATOM	2402	OG1	THR	A	16	-15.958	68.100	-6.758	1.00	29.56	T3
45	ATOM	2403	CG2	THR	A	16	-13.706	68.584	-7.442	1.00	31.77	T3
	ATOM	2404	C	THR	A	16	-16.644	70.741	-6.290	1.00	31.75	T3
	ATOM	2405	O	THR	A	16	-17.360	69.938	-5.690	1.00	29.57	T3
	ATOM	2406	N	ILE	A	17	-17.081	71.925	-6.703	1.00	32.59	T3
	ATOM	2407	CA	ILE	A	17	-18.450	72.364	-6.448	1.00	35.69	T3
50	ATOM	2408	CB	ILE	A	17	-18.720	73.696	-7.136	1.00	25.96	T3
	ATOM	2409	CG2	ILE	A	17	-20.146	74.131	-6.878	1.00	27.81	T3
	ATOM	2410	CG1	ILE	A	17	-17.745	74.750	-6.616	1.00	32.14	T3
	ATOM	2411	CD1	ILE	A	17	-17.868	76.099	-7.298	1.00	28.17	T3
	ATOM	2412	C	ILE	A	17	-19.551	71.378	-6.872	1.00	27.63	T3
55	ATOM	2413	O	ILE	A	17	-19.550	70.854	-8.000	1.00	30.08	T3
	ATOM	2414	N	GLN	A	18	-20.491	71.133	-5.952	1.00	33.38	T3
	ATOM	2415	CA	GLN	A	18	-21.615	70.230	-6.208	1.00	34.27	T3
	ATOM	2416	CB	GLN	A	18	-21.692	69.150	-5.141	1.00	30.08	T3
	ATOM	2417	CG	GLN	A	18	-21.672	67.768	-5.727	1.00	27.31	T3
60	ATOM	2418	CD	GLN	A	18	-20.303	67.409	-6.221	1.00	25.63	T3
	ATOM	2419	OE1	GLN	A	18	-19.410	67.098	-5.421	1.00	29.39	T3
	ATOM	2420	NE2	GLN	A	18	-20.105	67.471	-7.541	1.00	27.98	T3
	ATOM	2421	C	GLN	A	18	-22.938	70.993	-6.233	1.00	32.90	T3
	ATOM	2422	O	GLN	A	18	-23.207	71.823	-5.356	1.00	30.27	T3
65	ATOM	2423	N	LYS	A	19	-23.770	70.708	-7.231	1.00	34.96	T3
	ATOM	2424	CA	LYS	A	19	-25.046	71.401	-7.347	1.00	27.82	T3

	ATOM	2425	CB	LYS	A	19	-24.802	72.875	-7.645	1.00	34.58	T3
	ATOM	2426	CG	LYS	A	19	-26.052	73.637	-8.035	1.00	28.72	T3
	ATOM	2427	CD	LYS	A	19	-25.773	75.139	-8.094	1.00	30.32	T3
	ATOM	2428	CE	LYS	A	19	-27.020	75.930	-8.505	1.00	28.57	T3
5	ATOM	2429	NZ	LYS	A	19	-26.792	77.414	-8.533	1.00	28.12	T3
	ATOM	2430	C	LYS	A	19	-25.943	70.810	-8.424	1.00	34.16	T3
	ATOM	2431	O	LYS	A	19	-25.503	70.598	-9.559	1.00	26.60	T3
	ATOM	2432	N	GLY	A	20	-27.204	70.559	-8.068	1.00	30.57	T3
	ATOM	2433	CA	GLY	A	20	-28.151	69.981	-9.008	1.00	33.25	T3
10	ATOM	2434	C	GLY	A	20	-27.591	68.702	-9.599	1.00	26.54	T3
	ATOM	2435	O	GLY	A	20	-27.722	68.469	-10.805	1.00	24.97	T3
	ATOM	2436	N	SER	A	21	-26.965	67.881	-8.746	1.00	31.01	T3
	ATOM	2437	CA	SER	A	21	-26.347	66.610	-9.149	1.00	30.53	T3
	ATOM	2438	CB	SER	A	21	-27.425	65.558	-9.435	1.00	35.69	T3
15	ATOM	2439	OG	SER	A	21	-28.275	65.956	-10.494	1.00	29.83	T3
	ATOM	2440	C	SER	A	21	-25.412	66.769	-10.365	1.00	23.52	T3
	ATOM	2441	O	SER	A	21	-25.373	65.924	-11.272	1.00	31.14	T3
	ATOM	2442	N	TYR	A	22	-24.667	67.874	-10.358	1.00	25.30	T3
	ATOM	2443	CA	TYR	A	22	-23.704	68.207	-11.398	1.00	29.36	T3
20	ATOM	2444	CB	TYR	A	22	-24.161	69.441	-12.176	1.00	31.70	T3
	ATOM	2445	CG	TYR	A	22	-24.899	69.137	-13.446	1.00	30.36	T3
	ATOM	2446	CD1	TYR	A	22	-25.573	67.928	-13.614	1.00	25.19	T3
	ATOM	2447	CE1	TYR	A	22	-26.290	67.658	-14.778	1.00	30.60	T3
	ATOM	2448	CD2	TYR	A	22	-24.957	70.076	-14.475	1.00	29.24	T3
25	ATOM	2449	CE2	TYR	A	22	-25.676	69.820	-15.647	1.00	28.75	T3
	ATOM	2450	CZ	TYR	A	22	-26.341	68.604	-15.791	1.00	37.86	T3
	ATOM	2451	OH	TYR	A	22	-27.059	68.330	-16.941	1.00	27.34	T3
	ATOM	2452	C	TYR	A	22	-22.414	68.539	-10.674	1.00	30.33	T3
	ATOM	2453	O	TYR	A	22	-22.443	69.059	-9.553	1.00	30.80	T3
30	ATOM	2454	N	THR	A	23	-21.282	68.235	-11.294	1.00	34.73	T3
	ATOM	2455	CA	THR	A	23	-20.011	68.555	-10.671	1.00	28.46	T3
	ATOM	2456	CB	THR	A	23	-19.035	67.350	-10.680	1.00	34.95	T3
	ATOM	2457	OG1	THR	A	23	-19.742	66.140	-10.380	1.00	29.62	T3
	ATOM	2458	CG2	THR	A	23	-17.973	67.542	-9.613	1.00	25.24	T3
35	ATOM	2459	C	THR	A	23	-19.385	69.716	-11.438	1.00	30.01	T3
	ATOM	2460	O	THR	A	23	-19.313	69.702	-12.667	1.00	28.27	T3
	ATOM	2461	N	PHE	A	24	-18.950	70.730	-10.705	1.00	32.59	T3
	ATOM	2462	CA	PHE	A	24	-18.318	71.884	-11.321	1.00	35.49	T3
	ATOM	2463	CB	PHE	A	24	-19.112	73.144	-11.004	1.00	27.45	T3
40	ATOM	2464	CG	PHE	A	24	-20.478	73.160	-11.616	1.00	23.61	T3
	ATOM	2465	CD1	PHE	A	24	-21.530	72.470	-11.023	1.00	34.59	T3
	ATOM	2466	CD2	PHE	A	24	-20.712	73.850	-12.803	1.00	28.52	T3
	ATOM	2467	CE1	PHE	A	24	-22.797	72.469	-11.607	1.00	22.72	T3
	ATOM	2468	CE2	PHE	A	24	-21.969	73.851	-13.390	1.00	31.35	T3
45	ATOM	2469	CZ	PHE	A	24	-23.014	73.160	-12.791	1.00	29.96	T3
	ATOM	2470	C	PHE	A	24	-16.872	72.059	-10.864	1.00	33.43	T3
	ATOM	2471	O	PHE	A	24	-16.586	72.140	-9.662	1.00	27.08	T3
	ATOM	2472	N	VAL	A	25	-15.966	72.117	-11.836	1.00	31.58	T3
	ATOM	2473	CA	VAL	A	25	-14.549	72.283	-11.555	1.00	26.88	T3
50	ATOM	2474	CB	VAL	A	25	-13.714	72.163	-12.844	1.00	35.72	T3
	ATOM	2475	CG1	VAL	A	25	-12.250	72.412	-12.544	1.00	33.37	T3
	ATOM	2476	CG2	VAL	A	25	-13.896	70.792	-13.449	1.00	23.95	T3
	ATOM	2477	C	VAL	A	25	-14.274	73.643	-10.933	1.00	30.79	T3
	ATOM	2478	O	VAL	A	25	-14.782	74.656	-11.400	1.00	30.92	T3
55	ATOM	2479	N	PRO	A	26	-13.482	73.677	-9.851	1.00	39.28	T3
	ATOM	2480	CD	PRO	A	26	-13.007	72.513	-9.089	1.00	29.90	T3
	ATOM	2481	CA	PRO	A	26	-13.131	74.925	-9.164	1.00	25.89	T3
	ATOM	2482	CB	PRO	A	26	-12.546	74.446	-7.845	1.00	31.64	T3
	ATOM	2483	CG	PRO	A	26	-13.057	73.036	-7.702	1.00	30.88	T3
60	ATOM	2484	C	PRO	A	26	-12.062	75.611	-10.010	1.00	30.94	T3
	ATOM	2485	O	PRO	A	26	-10.935	75.123	-10.091	1.00	34.68	T3
	ATOM	2486	N	TRP	A	27	-12.394	76.732	-10.639	1.00	29.87	T3
	ATOM	2487	CA	TRP	A	27	-11.416	77.402	-11.486	1.00	37.98	T3
	ATOM	2488	CB	TRP	A	27	-12.120	78.140	-12.628	1.00	29.45	T3
65	ATOM	2489	CG	TRP	A	27	-12.853	77.230	-13.539	1.00	36.79	T3
	ATOM	2490	CD2	TRP	A	27	-12.319	76.092	-14.231	1.00	24.09	T3

	ATOM	2491	CE2	TRP	A	27	-13.385	75.495	-14.936	1.00	29.51	T3
	ATOM	2492	CE3	TRP	A	27	-11.045	75.517	-14.320	1.00	28.70	T3
	ATOM	2493	CD1	TRP	A	27	-14.178	77.281	-13.847	1.00	35.78	T3
	ATOM	2494	NE1	TRP	A	27	-14.508	76.239	-14.686	1.00	35.34	T3
5	ATOM	2495	CZ2	TRP	A	27	-13.217	74.352	-15.722	1.00	26.05	T3
	ATOM	2496	CZ3	TRP	A	27	-10.878	74.379	-15.101	1.00	34.77	T3
	ATOM	2497	CH2	TRP	A	27	-11.959	73.809	-15.791	1.00	37.02	T3
	ATOM	2498	C	TRP	A	27	-10.456	78.355	-10.788	1.00	34.47	T3
	ATOM	2499	O	TRP	A	27	-10.659	78.774	-9.648	1.00	38.45	T3
10	ATOM	2500	N	LEU	A	28	-9.393	78.680	-11.509	1.00	27.70	T3
	ATOM	2501	CA	LEU	A	28	-8.356	79.584	-11.042	1.00	32.63	T3
	ATOM	2502	CB	LEU	A	28	-7.267	78.794	-10.333	1.00	36.43	T3
	ATOM	2503	CG	LEU	A	28	-6.390	79.619	-9.406	1.00	32.44	T3
	ATOM	2504	CD1	LEU	A	28	-7.251	80.186	-8.272	1.00	34.56	T3
15	ATOM	2505	CD2	LEU	A	28	-5.273	78.737	-8.870	1.00	31.69	T3
	ATOM	2506	C	LEU	A	28	-7.793	80.254	-12.296	1.00	33.63	T3
	ATOM	2507	O	LEU	A	28	-7.432	79.579	-13.264	1.00	28.42	T3
	ATOM	2508	N	LEU	A	29	-7.728	81.578	-12.291	1.00	32.45	T3
	ATOM	2509	CA	LEU	A	29	-7.245	82.280	-13.459	1.00	30.92	T3
20	ATOM	2510	CB	LEU	A	29	-7.191	83.777	-13.196	1.00	22.68	T3
	ATOM	2511	CG	LEU	A	29	-6.649	84.561	-14.388	1.00	30.48	T3
	ATOM	2512	CD1	LEU	A	29	-7.713	84.644	-15.458	1.00	30.06	T3
	ATOM	2513	CD2	LEU	A	29	-6.223	85.939	-13.954	1.00	33.26	T3
	ATOM	2514	C	LEU	A	29	-5.875	81.818	-13.913	1.00	23.99	T3
25	ATOM	2515	O	LEU	A	29	-4.922	81.800	-13.136	1.00	29.58	T3
	ATOM	2516	N	SER	A	30	-5.779	81.433	-15.179	1.00	33.80	T3
	ATOM	2517	CA	SER	A	30	-4.500	81.029	-15.749	1.00	26.75	T3
	ATOM	2518	CB	SER	A	30	-4.700	80.072	-16.920	1.00	26.69	T3
	ATOM	2519	OG	SER	A	30	-3.466	79.783	-17.537	1.00	36.57	T3
30	ATOM	2520	C	SER	A	30	-3.891	82.330	-16.255	1.00	35.53	T3
	ATOM	2521	O	SER	A	30	-2.779	82.691	-15.894	1.00	32.14	T3
	ATOM	2522	N	PHE	A	31	-4.646	83.037	-17.087	1.00	30.60	T3
	ATOM	2523	CA	PHE	A	31	-4.209	84.313	-17.623	1.00	26.91	T3
	ATOM	2524	CB	PHE	A	31	-3.078	84.110	-18.636	1.00	33.44	T3
35	ATOM	2525	CG	PHE	A	31	-3.546	83.841	-20.037	1.00	27.42	T3
	ATOM	2526	CD1	PHE	A	31	-3.875	84.889	-20.885	1.00	27.99	T3
	ATOM	2527	CD2	PHE	A	31	-3.668	82.540	-20.504	1.00	35.21	T3
	ATOM	2528	CE1	PHE	A	31	-4.316	84.643	-22.171	1.00	31.33	T3
	ATOM	2529	CE2	PHE	A	31	-4.108	82.287	-21.789	1.00	25.12	T3
40	ATOM	2530	CZ	PHE	A	31	-4.433	83.338	-22.625	1.00	31.06	T3
	ATOM	2531	C	PHE	A	31	-5.398	85.005	-18.281	1.00	31.73	T3
	ATOM	2532	O	PHE	A	31	-6.347	84.354	-18.717	1.00	28.27	T3
	ATOM	2533	N	LYS	A	32	-5.352	86.328	-18.335	1.00	32.05	T3
	ATOM	2534	CA	LYS	A	32	-6.423	87.098	-18.944	1.00	29.14	T3
45	ATOM	2535	CB	LYS	A	32	-7.286	87.747	-17.872	1.00	24.48	T3
	ATOM	2536	CG	LYS	A	32	-8.211	88.785	-18.426	1.00	24.41	T3
	ATOM	2537	CD	LYS	A	32	-8.900	89.571	-17.347	1.00	28.51	T3
	ATOM	2538	CE	LYS	A	32	-9.594	90.762	-17.971	1.00	35.99	T3
	ATOM	2539	NZ	LYS	A	32	-10.432	91.480	-16.981	1.00	30.41	T3
50	ATOM	2540	C	LYS	A	32	-5.800	88.171	-19.821	1.00	29.14	T3
	ATOM	2541	O	LYS	A	32	-4.962	88.944	-19.365	1.00	31.41	T3
	ATOM	2542	N	ARG	A	33	-6.213	88.219	-21.080	1.00	30.09	T3
	ATOM	2543	CA	ARG	A	33	-5.667	89.179	-22.027	1.00	37.72	T3
	ATOM	2544	CB	ARG	A	33	-4.822	88.428	-23.054	1.00	26.20	T3
55	ATOM	2545	CG	ARG	A	33	-4.299	89.232	-24.210	1.00	24.58	T3
	ATOM	2546	CD	ARG	A	33	-3.107	88.497	-24.828	1.00	36.55	T3
	ATOM	2547	NE	ARG	A	33	-2.577	89.155	-26.025	1.00	31.34	T3
	ATOM	2548	CZ	ARG	A	33	-3.095	89.011	-27.241	1.00	35.36	T3
	ATOM	2549	NH1	ARG	A	33	-4.154	88.224	-27.419	1.00	36.78	T3
60	ATOM	2550	NH2	ARG	A	33	-2.569	89.663	-28.269	1.00	34.72	T3
	ATOM	2551	C	ARG	A	33	-6.783	89.936	-22.707	1.00	30.63	T3
	ATOM	2552	O	ARG	A	33	-7.670	89.340	-23.313	1.00	24.50	T3
	ATOM	2553	N	GLY	A	34	-6.750	91.254	-22.591	1.00	31.10	T3
	ATOM	2554	CA	GLY	A	34	-7.778	92.060	-23.215	1.00	27.96	T3
65	ATOM	2555	C	GLY	A	34	-8.926	92.397	-22.290	1.00	31.68	T3
	ATOM	2556	O	GLY	A	34	-8.831	92.227	-21.072	1.00	32.28	T3

	ATOM	2557	N	SER	A	35	-10.028	92.855	-22.877	1.00	28.76	T3
	ATOM	2558	CA	SER	A	35	-11.200	93.249	-22.107	1.00	24.95	T3
	ATOM	2559	CB	SER	A	35	-11.471	94.728	-22.339	1.00	27.96	T3
	ATOM	2560	OG	SER	A	35	-11.597	94.991	-23.728	1.00	30.14	T3
5	ATOM	2561	C	SER	A	35	-12.485	92.471	-22.390	1.00	29.40	T3
	ATOM	2562	O	SER	A	35	-13.386	92.463	-21.551	1.00	29.27	T3
	ATOM	2563	N	ALA	A	36	-12.572	91.821	-23.550	1.00	29.27	T3
	ATOM	2564	CA	ALA	A	36	-13.779	91.096	-23.930	1.00	30.92	T3
	ATOM	2565	CB	ALA	A	36	-13.652	90.600	-25.358	1.00	34.42	T3
10	ATOM	2566	C	ALA	A	36	-14.214	89.947	-23.029	1.00	37.51	T3
	ATOM	2567	O	ALA	A	36	-15.365	89.511	-23.108	1.00	28.88	T3
	ATOM	2568	N	LEU	A	37	-13.320	89.454	-22.173	1.00	30.61	T3
	ATOM	2569	CA	LEU	A	37	-13.668	88.338	-21.288	1.00	30.66	T3
	ATOM	2570	CB	LEU	A	37	-13.089	87.041	-21.847	1.00	35.92	T3
15	ATOM	2571	CG	LEU	A	37	-13.599	86.671	-23.240	1.00	28.56	T3
	ATOM	2572	CD1	LEU	A	37	-12.675	85.674	-23.888	1.00	29.60	T3
	ATOM	2573	CD2	LEU	A	37	-15.004	86.129	-23.134	1.00	30.66	T3
	ATOM	2574	C	LEU	A	37	-13.193	88.540	-19.856	1.00	30.08	T3
	ATOM	2575	O	LEU	A	37	-12.154	89.149	-19.622	1.00	30.97	T3
20	ATOM	2576	N	GLU	A	38	-13.960	88.016	-18.902	1.00	29.95	T3
	ATOM	2577	CA	GLU	A	38	-13.651	88.148	-17.472	1.00	30.49	T3
	ATOM	2578	CB	GLU	A	38	-14.394	89.348	-16.880	1.00	32.77	T3
	ATOM	2579	CG	GLU	A	38	-13.738	90.696	-17.083	1.00	34.01	T3
	ATOM	2580	CD	GLU	A	38	-14.658	91.847	-16.697	1.00	32.52	T3
25	ATOM	2581	OE1	GLU	A	38	-15.442	91.692	-15.726	1.00	32.14	T3
	ATOM	2582	OE2	GLU	A	38	-14.589	92.910	-17.360	1.00	30.64	T3
	ATOM	2583	C	GLU	A	38	-14.071	86.923	-16.673	1.00	29.78	T3
	ATOM	2584	O	GLU	A	38	-14.855	86.103	-17.152	1.00	28.82	T3
	ATOM	2585	N	GLU	A	39	-13.549	86.799	-15.452	1.00	29.36	T3
30	ATOM	2586	CA	GLU	A	39	-13.934	85.688	-14.580	1.00	27.84	T3
	ATOM	2587	CB	GLU	A	39	-12.856	85.327	-13.582	1.00	29.77	T3
	ATOM	2588	CG	GLU	A	39	-11.476	85.283	-14.106	1.00	26.80	T3
	ATOM	2589	CD	GLU	A	39	-10.477	85.622	-13.010	1.00	28.12	T3
	ATOM	2590	OE1	GLU	A	39	-10.060	86.810	-12.936	1.00	32.42	T3
35	ATOM	2591	OE2	GLU	A	39	-10.132	84.708	-12.213	1.00	26.13	T3
	ATOM	2592	C	GLU	A	39	-15.077	86.219	-13.755	1.00	34.72	T3
	ATOM	2593	O	GLU	A	39	-15.102	87.398	-13.409	1.00	35.32	T3
	ATOM	2594	N	LYS	A	40	-16.017	85.357	-13.419	1.00	35.71	T3
	ATOM	2595	CA	LYS	A	40	-17.124	85.790	-12.596	1.00	30.70	T3
40	ATOM	2596	CB	LYS	A	40	-18.142	86.584	-13.409	1.00	21.51	T3
	ATOM	2597	CG	LYS	A	40	-19.301	87.096	-12.559	1.00	25.80	T3
	ATOM	2598	CD	LYS	A	40	-20.478	87.524	-13.423	1.00	27.10	T3
	ATOM	2599	CE	LYS	A	40	-21.708	87.835	-12.571	1.00	32.20	T3
	ATOM	2600	NZ	LYS	A	40	-22.904	88.094	-13.442	1.00	26.87	T3
45	ATOM	2601	C	LYS	A	40	-17.791	84.586	-11.967	1.00	30.48	T3
	ATOM	2602	O	LYS	A	40	-18.511	83.828	-12.621	1.00	32.70	T3
	ATOM	2603	N	GLU	A	41	-17.519	84.400	-10.686	1.00	30.46	T3
	ATOM	2604	CA	GLU	A	41	-18.105	83.298	-9.946	1.00	27.57	T3
	ATOM	2605	CB	GLU	A	41	-19.574	83.613	-9.682	1.00	27.35	T3
50	ATOM	2606	CG	GLU	A	41	-19.743	85.001	-9.074	1.00	33.62	T3
	ATOM	2607	CD	GLU	A	41	-21.175	85.508	-9.142	1.00	30.94	T3
	ATOM	2608	OE1	GLU	A	41	-21.773	85.501	-10.257	1.00	25.39	T3
	ATOM	2609	OE2	GLU	A	41	-21.695	85.927	-8.077	1.00	28.91	T3
	ATOM	2610	C	GLU	A	41	-17.950	81.974	-10.685	1.00	35.14	T3
55	ATOM	2611	O	GLU	A	41	-18.929	81.317	-11.036	1.00	31.41	T3
	ATOM	2612	N	ASN	A	42	-16.698	81.612	-10.932	1.00	31.27	T3
	ATOM	2613	CA	ASN	A	42	-16.364	80.365	-11.585	1.00	28.52	T3
	ATOM	2614	CB	ASN	A	42	-16.849	79.199	-10.742	1.00	19.92	T3
	ATOM	2615	CG	ASN	A	42	-15.830	78.092	-10.660	1.00	27.58	T3
60	ATOM	2616	OD1	ASN	A	42	-16.132	76.931	-10.949	1.00	29.50	T3
	ATOM	2617	ND2	ASN	A	42	-14.605	78.443	-10.260	1.00	28.77	T3
	ATOM	2618	C	ASN	A	42	-16.891	80.203	-12.993	1.00	32.01	T3
	ATOM	2619	O	ASN	A	42	-16.992	79.083	-13.492	1.00	36.16	T3
	ATOM	2620	N	LYS	A	43	-17.225	81.317	-13.630	1.00	29.12	T3
65	ATOM	2621	CA	LYS	A	43	-17.722	81.289	-14.996	1.00	31.74	T3
	ATOM	2622	CB	LYS	A	43	-19.225	81.557	-15.021	1.00	23.80	T3

	ATOM	2623	CG	LYS	A	43	-20.057	80.439	-14.431	1.00	28.23	T3
	ATOM	2624	CD	LYS	A	43	-21.535	80.789	-14.419	1.00	27.42	T3
	ATOM	2625	CE	LYS	A	43	-21.836	81.855	-13.383	1.00	26.04	T3
	ATOM	2626	NZ	LYS	A	43	-23.293	82.203	-13.332	1.00	29.48	T3
5	ATOM	2627	C	LYS	A	43	-17.006	82.354	-15.805	1.00	32.91	T3
	ATOM	2628	O	LYS	A	43	-16.414	83.273	-15.242	1.00	38.09	T3
	ATOM	2629	N	ILE	A	44	-17.044	82.229	-17.124	1.00	32.23	T3
	ATOM	2630	CA	ILE	A	44	-16.413	83.223	-17.968	1.00	32.82	T3
	ATOM	2631	CB	ILE	A	44	-15.752	82.584	-19.186	1.00	28.14	T3
10	ATOM	2632	CG2	ILE	A	44	-15.111	83.655	-20.039	1.00	29.04	T3
	ATOM	2633	CG1	ILE	A	44	-14.707	81.573	-18.732	1.00	26.10	T3
	ATOM	2634	CD1	ILE	A	44	-14.008	80.872	-19.865	1.00	33.85	T3
	ATOM	2635	C	ILE	A	44	-17.496	84.184	-18.432	1.00	27.18	T3
	ATOM	2636	O	ILE	A	44	-18.500	83.778	-19.014	1.00	31.18	T3
15	ATOM	2637	N	LEU	A	45	-17.288	85.463	-18.165	1.00	26.05	T3
	ATOM	2638	CA	LEU	A	45	-18.252	86.485	-18.538	1.00	32.26	T3
	ATOM	2639	CB	LEU	A	45	-18.368	87.504	-17.411	1.00	25.43	T3
	ATOM	2640	CG	LEU	A	45	-19.308	88.666	-17.725	1.00	35.22	T3
	ATOM	2641	CD1	LEU	A	45	-20.760	88.176	-17.764	1.00	24.04	T3
20	ATOM	2642	CD2	LEU	A	45	-19.131	89.737	-16.671	1.00	33.01	T3
	ATOM	2643	C	LEU	A	45	-17.898	87.210	-19.832	1.00	31.52	T3
	ATOM	2644	O	LEU	A	45	-16.782	87.709	-19.987	1.00	33.87	T3
	ATOM	2645	N	VAL	A	46	-18.857	87.286	-20.750	1.00	30.84	T3
	ATOM	2646	CA	VAL	A	46	-18.635	87.956	-22.028	1.00	27.13	T3
25	ATOM	2647	CB	VAL	A	46	-19.556	87.385	-23.114	1.00	29.90	T3
	ATOM	2648	CG1	VAL	A	46	-19.316	88.101	-24.418	1.00	29.41	T3
	ATOM	2649	CG2	VAL	A	46	-19.301	85.909	-23.279	1.00	32.61	T3
	ATOM	2650	C	VAL	A	46	-18.886	89.457	-21.911	1.00	32.91	T3
	ATOM	2651	O	VAL	A	46	-19.989	89.877	-21.584	1.00	29.93	T3
30	ATOM	2652	N	LYS	A	47	-17.868	90.266	-22.184	1.00	35.78	T3
	ATOM	2653	CA	LYS	A	47	-18.012	91.716	-22.081	1.00	32.26	T3
	ATOM	2654	CB	LYS	A	47	-16.810	92.313	-21.352	1.00	34.34	T3
	ATOM	2655	CG	LYS	A	47	-16.823	92.095	-19.856	1.00	28.95	T3
	ATOM	2656	CD	LYS	A	47	-18.122	92.610	-19.269	1.00	32.76	T3
35	ATOM	2657	CE	LYS	A	47	-18.074	92.710	-17.752	1.00	30.78	T3
	ATOM	2658	NZ	LYS	A	47	-17.245	93.862	-17.290	1.00	31.04	T3
	ATOM	2659	C	LYS	A	47	-18.202	92.446	-23.408	1.00	36.58	T3
	ATOM	2660	O	LYS	A	47	-18.483	93.639	-23.427	1.00	30.56	T3
	ATOM	2661	N	GLU	A	48	-18.037	91.730	-24.514	1.00	30.18	T3
40	ATOM	2662	CA	GLU	A	48	-18.191	92.304	-25.848	1.00	32.85	T3
	ATOM	2663	CB	GLU	A	48	-16.848	92.684	-26.441	1.00	29.87	T3
	ATOM	2664	CG	GLU	A	48	-16.173	93.852	-25.802	1.00	29.58	T3
	ATOM	2665	CD	GLU	A	48	-14.782	94.059	-26.356	1.00	34.92	T3
	ATOM	2666	OE1	GLU	A	48	-14.610	93.932	-27.596	1.00	32.36	T3
45	ATOM	2667	OE2	GLU	A	48	-13.870	94.351	-25.544	1.00	24.64	T3
	ATOM	2668	C	GLU	A	48	-18.769	91.229	-26.724	1.00	26.43	T3
	ATOM	2669	O	GLU	A	48	-18.201	90.138	-26.816	1.00	34.10	T3
	ATOM	2670	N	THR	A	49	-19.884	91.518	-27.386	1.00	22.49	T3
	ATOM	2671	CA	THR	A	49	-20.481	90.499	-28.237	1.00	29.42	T3
50	ATOM	2672	CB	THR	A	49	-21.892	90.895	-28.683	1.00	29.83	T3
	ATOM	2673	OG1	THR	A	49	-21.813	91.591	-29.921	1.00	26.63	T3
	ATOM	2674	CG2	THR	A	49	-22.543	91.798	-27.649	1.00	30.47	T3
	ATOM	2675	C	THR	A	49	-19.580	90.276	-29.454	1.00	24.80	T3
	ATOM	2676	O	THR	A	49	-18.874	91.186	-29.896	1.00	27.68	T3
55	ATOM	2677	N	GLY	A	50	-19.591	89.053	-29.968	1.00	33.54	T3
	ATOM	2678	CA	GLY	A	50	-18.773	88.727	-31.116	1.00	35.58	T3
	ATOM	2679	C	GLY	A	50	-18.661	87.227	-31.267	1.00	24.87	T3
	ATOM	2680	O	GLY	A	50	-19.433	86.489	-30.664	1.00	35.49	T3
	ATOM	2681	N	TYR	A	51	-17.707	86.771	-32.074	1.00	30.25	T3
60	ATOM	2682	CA	TYR	A	51	-17.492	85.339	-32.291	1.00	29.64	T3
	ATOM	2683	CB	TYR	A	51	-17.259	85.058	-33.777	1.00	21.90	T3
	ATOM	2684	CG	TYR	A	51	-18.529	85.147	-34.584	1.00	28.96	T3
	ATOM	2685	CD1	TYR	A	51	-19.056	86.378	-34.962	1.00	35.75	T3
	ATOM	2686	CE1	TYR	A	51	-20.290	86.461	-35.595	1.00	32.73	T3
65	ATOM	2687	CD2	TYR	A	51	-19.265	83.996	-34.873	1.00	30.59	T3
	ATOM	2688	CE2	TYR	A	51	-20.496	84.066	-35.502	1.00	32.04	T3

	ATOM	2689	CZ	TYR	A	51	-21.009	85.296	-35.854	1.00	30.70	T3
	ATOM	2690	OH	TYR	A	51	-22.273	85.354	-36.405	1.00	22.84	T3
	ATOM	2691	C	TYR	A	51	-16.306	84.842	-31.470	1.00	27.19	T3
	ATOM	2692	O	TYR	A	51	-15.219	85.411	-31.521	1.00	33.12	T3
5	ATOM	2693	N	PHE	A	52	-16.520	83.779	-30.708	1.00	31.24	T3
	ATOM	2694	CA	PHE	A	52	-15.465	83.247	-29.867	1.00	38.20	T3
	ATOM	2695	CB	PHE	A	52	-15.817	83.424	-28.386	1.00	26.46	T3
	ATOM	2696	CG	PHE	A	52	-16.061	84.844	-27.973	1.00	35.21	T3
	ATOM	2697	CD1	PHE	A	52	-17.236	85.494	-28.324	1.00	26.29	T3
10	ATOM	2698	CD2	PHE	A	52	-15.120	85.527	-27.219	1.00	34.48	T3
	ATOM	2699	CE1	PHE	A	52	-17.470	86.805	-27.930	1.00	27.17	T3
	ATOM	2700	CE2	PHE	A	52	-15.341	86.836	-26.818	1.00	30.56	T3
	ATOM	2701	CZ	PHE	A	52	-16.518	87.479	-27.174	1.00	26.20	T3
	ATOM	2702	C	PHE	A	52	-15.178	81.776	-30.105	1.00	30.39	T3
15	ATOM	2703	O	PHE	A	52	-16.065	81.002	-30.457	1.00	26.62	T3
	ATOM	2704	N	PHE	A	53	-13.914	81.411	-29.910	1.00	33.22	T3
	ATOM	2705	CA	PHE	A	53	-13.460	80.034	-30.021	1.00	31.95	T3
	ATOM	2706	CB	PHE	A	53	-12.051	79.975	-30.585	1.00	30.17	T3
	ATOM	2707	CG	PHE	A	53	-11.440	78.613	-30.530	1.00	27.76	T3
20	ATOM	2708	CD1	PHE	A	53	-11.937	77.582	-31.315	1.00	34.02	T3
	ATOM	2709	CD2	PHE	A	53	-10.373	78.353	-29.675	1.00	33.66	T3
	ATOM	2710	CE1	PHE	A	53	-11.383	76.310	-31.252	1.00	27.79	T3
	ATOM	2711	CE2	PHE	A	53	-9.809	77.088	-29.601	1.00	31.93	T3
	ATOM	2712	CZ	PHE	A	53	-10.313	76.062	-30.389	1.00	23.06	T3
25	ATOM	2713	C	PHE	A	53	-13.437	79.598	-28.570	1.00	30.80	T3
	ATOM	2714	O	PHE	A	53	-12.780	80.237	-27.754	1.00	27.34	T3
	ATOM	2715	N	ILE	A	54	-14.155	78.530	-28.243	1.00	32.32	T3
	ATOM	2716	CA	ILE	A	54	-14.220	78.066	-26.865	1.00	28.78	T3
	ATOM	2717	CB	ILE	A	54	-15.657	78.149	-26.355	1.00	30.15	T3
30	ATOM	2718	CG2	ILE	A	54	-15.698	77.877	-24.868	1.00	33.26	T3
	ATOM	2719	CG1	ILE	A	54	-16.210	79.539	-26.658	1.00	26.98	T3
	ATOM	2720	CD1	ILE	A	54	-17.689	79.625	-26.622	1.00	31.67	T3
	ATOM	2721	C	ILE	A	54	-13.715	76.644	-26.745	1.00	33.11	T3
	ATOM	2722	O	ILE	A	54	-14.122	75.778	-27.510	1.00	36.44	T3
35	ATOM	2723	N	TYR	A	55	-12.826	76.409	-25.783	1.00	31.24	T3
	ATOM	2724	CA	TYR	A	55	-12.257	75.080	-25.587	1.00	23.11	T3
	ATOM	2725	CB	TYR	A	55	-10.824	75.044	-26.119	1.00	32.98	T3
	ATOM	2726	CG	TYR	A	55	-9.914	76.070	-25.493	1.00	30.21	T3
	ATOM	2727	CD1	TYR	A	55	-9.098	75.744	-24.414	1.00	26.57	T3
40	ATOM	2728	CE1	TYR	A	55	-8.278	76.694	-23.817	1.00	36.80	T3
	ATOM	2729	CD2	TYR	A	55	-9.887	77.378	-25.961	1.00	28.70	T3
	ATOM	2730	CE2	TYR	A	55	-9.075	78.337	-25.371	1.00	32.96	T3
	ATOM	2731	CZ	TYR	A	55	-8.275	77.989	-24.300	1.00	29.90	T3
	ATOM	2732	OH	TYR	A	55	-7.488	78.942	-23.705	1.00	25.20	T3
45	ATOM	2733	C	TYR	A	55	-12.281	74.648	-24.133	1.00	36.33	T3
	ATOM	2734	O	TYR	A	55	-12.593	75.434	-23.249	1.00	32.43	T3
	ATOM	2735	N	GLY	A	56	-11.957	73.387	-23.890	1.00	28.09	T3
	ATOM	2736	CA	GLY	A	56	-11.951	72.891	-22.531	1.00	35.30	T3
	ATOM	2737	C	GLY	A	56	-11.598	71.423	-22.474	1.00	31.20	T3
50	ATOM	2738	O	GLY	A	56	-12.033	70.643	-23.319	1.00	36.84	T3
	ATOM	2739	N	GLN	A	57	-10.789	71.055	-21.485	1.00	35.01	T3
	ATOM	2740	CA	GLN	A	57	-10.375	69.671	-21.285	1.00	32.02	T3
	ATOM	2741	CB	GLN	A	57	-8.924	69.461	-21.733	1.00	30.74	T3
	ATOM	2742	CG	GLN	A	57	-8.380	68.067	-21.396	1.00	21.27	T3
55	ATOM	2743	CD	GLN	A	57	-6.991	67.802	-21.944	1.00	35.75	T3
	ATOM	2744	OE1	GLN	A	57	-6.807	67.656	-23.144	1.00	26.80	T3
	ATOM	2745	NE2	GLN	A	57	-6.009	67.735	-21.063	1.00	28.12	T3
	ATOM	2746	C	GLN	A	57	-10.507	69.285	-19.813	1.00	27.58	T3
	ATOM	2747	O	GLN	A	57	-10.330	70.117	-18.926	1.00	33.27	T3
60	ATOM	2748	N	VAL	A	58	-10.825	68.017	-19.571	1.00	32.50	T3
	ATOM	2749	CA	VAL	A	58	-10.977	67.488	-18.220	1.00	36.09	T3
	ATOM	2750	CB	VAL	A	58	-12.470	67.371	-17.819	1.00	33.17	T3
	ATOM	2751	CG1	VAL	A	58	-12.609	66.569	-16.542	1.00	34.47	T3
	ATOM	2752	CG2	VAL	A	58	-13.066	68.745	-17.633	1.00	31.24	T3
65	ATOM	2753	C	VAL	A	58	-10.364	66.094	-18.184	1.00	23.11	T3
	ATOM	2754	O	VAL	A	58	-10.502	65.343	-19.144	1.00	30.32	T3

	ATOM	2755	N	LEU	A	59	-9.676	65.756	-17.095	1.00	38.18	T3
	ATOM	2756	CA	LEU	A	59	-9.074	64.433	-16.951	1.00	31.95	T3
	ATOM	2757	CB	LEU	A	59	-7.679	64.534	-16.328	1.00	33.05	T3
	ATOM	2758	CG	LEU	A	59	-6.715	63.374	-16.610	1.00	32.19	T3
5	ATOM	2759	CD1	LEU	A	59	-5.598	63.400	-15.595	1.00	34.42	T3
	ATOM	2760	CD2	LEU	A	59	-7.423	62.046	-16.532	1.00	26.57	T3
	ATOM	2761	C	LEU	A	59	-9.982	63.603	-16.035	1.00	37.68	T3
	ATOM	2762	O	LEU	A	59	-10.126	63.902	-14.843	1.00	26.27	T3
	ATOM	2763	N	TYR	A	60	-10.593	62.560	-16.587	1.00	31.84	T3
10	ATOM	2764	CA	TYR	A	60	-11.480	61.715	-15.797	1.00	29.53	T3
	ATOM	2765	CB	TYR	A	60	-12.633	61.217	-16.659	1.00	23.72	T3
	ATOM	2766	CG	TYR	A	60	-13.440	62.347	-17.209	1.00	25.89	T3
	ATOM	2767	CD1	TYR	A	60	-13.339	62.711	-18.546	1.00	31.30	T3
	ATOM	2768	CE1	TYR	A	60	-14.029	63.810	-19.040	1.00	22.26	T3
15	ATOM	2769	CD2	TYR	A	60	-14.255	63.103	-16.375	1.00	33.94	T3
	ATOM	2770	CE2	TYR	A	60	-14.941	64.195	-16.849	1.00	34.04	T3
	ATOM	2771	CZ	TYR	A	60	-14.824	64.548	-18.184	1.00	33.94	T3
	ATOM	2772	OH	TYR	A	60	-15.488	65.653	-18.658	1.00	28.99	T3
	ATOM	2773	C	TYR	A	60	-10.790	60.523	-15.157	1.00	26.32	T3
20	ATOM	2774	O	TYR	A	60	-10.131	59.730	-15.834	1.00	28.85	T3
	ATOM	2775	N	THR	A	61	-10.949	60.398	-13.846	1.00	41.95	T3
	ATOM	2776	CA	THR	A	61	-10.354	59.291	-13.120	1.00	31.98	T3
	ATOM	2777	CB	THR	A	61	-9.378	59.796	-12.068	1.00	34.33	T3
	ATOM	2778	OG1	THR	A	61	-10.032	60.761	-11.234	1.00	36.41	T3
25	ATOM	2779	CG2	THR	A	61	-8.181	60.438	-12.739	1.00	29.55	T3
	ATOM	2780	C	THR	A	61	-11.468	58.498	-12.458	1.00	34.53	T3
	ATOM	2781	O	THR	A	61	-11.234	57.685	-11.572	1.00	31.87	T3
	ATOM	2782	N	ASP	A	62	-12.688	58.754	-12.911	1.00	33.54	T3
	ATOM	2783	CA	ASP	A	62	-13.883	58.091	-12.409	1.00	29.43	T3
30	ATOM	2784	CB	ASP	A	62	-15.085	59.016	-12.635	1.00	28.54	T3
	ATOM	2785	CG	ASP	A	62	-16.317	58.573	-11.883	1.00	33.22	T3
	ATOM	2786	OD1	ASP	A	62	-16.918	59.431	-11.185	1.00	27.03	T3
	ATOM	2787	OD2	ASP	A	62	-16.682	57.375	-11.998	1.00	27.47	T3
	ATOM	2788	C	ASP	A	62	-14.031	56.791	-13.201	1.00	27.14	T3
35	ATOM	2789	O	ASP	A	62	-13.632	56.721	-14.360	1.00	23.06	T3
	ATOM	2790	N	LYS	A	63	-14.595	55.754	-12.598	1.00	34.63	T3
	ATOM	2791	CA	LYS	A	63	-14.732	54.501	-13.334	1.00	30.46	T3
	ATOM	2792	CB	LYS	A	63	-14.152	53.343	-12.521	1.00	32.20	T3
	ATOM	2793	CG	LYS	A	63	-14.826	53.137	-11.171	1.00	34.88	T3
40	ATOM	2794	CD	LYS	A	63	-14.219	51.939	-10.428	1.00	32.49	T3
	ATOM	2795	CE	LYS	A	63	-12.717	52.135	-10.141	1.00	35.92	T3
	ATOM	2796	NZ	LYS	A	63	-12.100	50.964	-9.429	1.00	31.69	T3
	ATOM	2797	C	LYS	A	63	-16.162	54.160	-13.736	1.00	29.66	T3
	ATOM	2798	O	LYS	A	63	-16.475	52.992	-14.003	1.00	30.55	T3
45	ATOM	2799	N	THR	A	64	-17.029	55.162	-13.811	1.00	24.28	T3
	ATOM	2800	CA	THR	A	64	-18.412	54.866	-14.165	1.00	32.10	T3
	ATOM	2801	CB	THR	A	64	-19.400	55.876	-13.502	1.00	36.12	T3
	ATOM	2802	OG1	THR	A	64	-18.985	57.219	-13.769	1.00	30.30	T3
	ATOM	2803	CG2	THR	A	64	-19.452	55.649	-11.991	1.00	36.31	T3
50	ATOM	2804	C	THR	A	64	-18.733	54.739	-15.659	1.00	22.67	T3
	ATOM	2805	O	THR	A	64	-19.554	55.482	-16.193	1.00	33.95	T3
	ATOM	2806	N	TYR	A	65	-18.078	53.781	-16.314	1.00	25.25	T3
	ATOM	2807	CA	TYR	A	65	-18.290	53.470	-17.734	1.00	30.14	T3
	ATOM	2808	CB	TYR	A	65	-19.550	52.601	-17.875	1.00	29.99	T3
55	ATOM	2809	CG	TYR	A	65	-20.777	53.337	-18.370	1.00	32.92	T3
	ATOM	2810	CD1	TYR	A	65	-21.084	53.392	-19.736	1.00	26.09	T3
	ATOM	2811	CE1	TYR	A	65	-22.217	54.079	-20.201	1.00	27.84	T3
	ATOM	2812	CD2	TYR	A	65	-21.630	53.985	-17.477	1.00	28.16	T3
	ATOM	2813	CE2	TYR	A	65	-22.763	54.675	-17.926	1.00	27.27	T3
60	ATOM	2814	CZ	TYR	A	65	-23.052	54.720	-19.285	1.00	29.14	T3
	ATOM	2815	OH	TYR	A	65	-24.174	55.406	-19.713	1.00	29.46	T3
	ATOM	2816	C	TYR	A	65	-18.352	54.609	-18.764	1.00	24.97	T3
	ATOM	2817	O	TYR	A	65	-18.087	54.384	-19.948	1.00	34.47	T3
	ATOM	2818	N	ALA	A	66	-18.703	55.818	-18.344	1.00	30.34	T3
65	ATOM	2819	CA	ALA	A	66	-18.782	56.921	-19.291	1.00	25.58	T3
	ATOM	2820	CB	ALA	A	66	-20.044	56.785	-20.131	1.00	39.12	T3

	ATOM	2821	C	ALA	A	66	-18.764	58.273	-18.590	1.00	34.07	T3
	ATOM	2822	O	ALA	A	66	-19.662	58.589	-17.803	1.00	27.28	T3
	ATOM	2823	N	MET	A	67	-17.739	59.069	-18.875	1.00	37.49	T3
	ATOM	2824	CA	MET	A	67	-17.626	60.390	-18.282	1.00	31.58	T3
5	ATOM	2825	CB	MET	A	67	-16.353	60.487	-17.443	1.00	35.21	T3
	ATOM	2826	CG	MET	A	67	-16.393	59.663	-16.167	1.00	28.28	T3
	ATOM	2827	SD	MET	A	67	-17.759	60.182	-15.079	1.00	23.37	T3
	ATOM	2828	CE	MET	A	67	-17.062	61.631	-14.289	1.00	21.72	T3
	ATOM	2829	C	MET	A	67	-17.596	61.423	-19.389	1.00	23.01	T3
10	ATOM	2830	O	MET	A	67	-17.451	61.075	-20.564	1.00	28.47	T3
	ATOM	2831	N	GLY	A	68	-17.741	62.692	-19.017	1.00	29.23	T3
	ATOM	2832	CA	GLY	A	68	-17.719	63.759	-20.004	1.00	29.25	T3
	ATOM	2833	C	GLY	A	68	-18.154	65.090	-19.428	1.00	29.38	T3
	ATOM	2834	O	GLY	A	68	-18.638	65.154	-18.292	1.00	38.85	T3
15	ATOM	2835	N	HIS	A	69	-17.969	66.163	-20.194	1.00	32.09	T3
	ATOM	2836	CA	HIS	A	69	-18.371	67.487	-19.736	1.00	28.09	T3
	ATOM	2837	CB	HIS	A	69	-17.167	68.287	-19.218	1.00	29.83	T3
	ATOM	2838	CG	HIS	A	69	-16.032	68.392	-20.187	1.00	27.72	T3
	ATOM	2839	CD2	HIS	A	69	-15.614	69.423	-20.956	1.00	33.53	T3
20	ATOM	2840	ND1	HIS	A	69	-15.151	67.359	-20.422	1.00	25.94	T3
	ATOM	2841	CE1	HIS	A	69	-14.236	67.748	-21.290	1.00	29.90	T3
	ATOM	2842	NE2	HIS	A	69	-14.495	68.997	-21.630	1.00	31.54	T3
	ATOM	2843	C	HIS	A	69	-19.100	68.281	-20.809	1.00	33.49	T3
	ATOM	2844	O	HIS	A	69	-19.114	67.905	-21.982	1.00	36.06	T3
25	ATOM	2845	N	LEU	A	70	-19.713	69.378	-20.384	1.00	28.31	T3
	ATOM	2846	CA	LEU	A	70	-20.465	70.253	-21.275	1.00	22.39	T3
	ATOM	2847	CB	LEU	A	70	-21.932	70.283	-20.868	1.00	27.17	T3
	ATOM	2848	CG	LEU	A	70	-22.577	68.981	-20.417	1.00	30.21	T3
	ATOM	2849	CD1	LEU	A	70	-23.948	69.292	-19.857	1.00	24.32	T3
30	ATOM	2850	CD2	LEU	A	70	-22.668	68.015	-21.575	1.00	31.33	T3
	ATOM	2851	C	LEU	A	70	-19.954	71.683	-21.191	1.00	27.29	T3
	ATOM	2852	O	LEU	A	70	-19.563	72.151	-20.125	1.00	28.88	T3
	ATOM	2853	N	ILE	A	71	-19.957	72.379	-22.317	1.00	29.13	T3
	ATOM	2854	CA	ILE	A	71	-19.558	73.773	-22.318	1.00	33.11	T3
35	ATOM	2855	CB	ILE	A	71	-18.547	74.069	-23.427	1.00	37.48	T3
	ATOM	2856	CG2	ILE	A	71	-18.399	75.564	-23.605	1.00	33.35	T3
	ATOM	2857	CG1	ILE	A	71	-17.198	73.441	-23.068	1.00	28.87	T3
	ATOM	2858	CD1	ILE	A	71	-16.129	73.657	-24.116	1.00	29.51	T3
	ATOM	2859	C	ILE	A	71	-20.873	74.487	-22.575	1.00	29.84	T3
40	ATOM	2860	O	ILE	A	71	-21.432	74.406	-23.666	1.00	33.65	T3
	ATOM	2861	N	GLN	A	72	-21.385	75.176	-21.566	1.00	29.23	T3
	ATOM	2862	CA	GLN	A	72	-22.673	75.840	-21.716	1.00	26.03	T3
	ATOM	2863	CB	GLN	A	72	-23.594	75.376	-20.603	1.00	28.49	T3
	ATOM	2864	CG	GLN	A	72	-23.536	73.884	-20.406	1.00	35.51	T3
45	ATOM	2865	CD	GLN	A	72	-24.587	73.409	-19.447	1.00	23.06	T3
	ATOM	2866	OE1	GLN	A	72	-24.685	73.899	-18.312	1.00	33.08	T3
	ATOM	2867	NE2	GLN	A	72	-25.395	72.447	-19.893	1.00	27.78	T3
	ATOM	2868	C	GLN	A	72	-22.673	77.358	-21.770	1.00	27.44	T3
	ATOM	2869	O	GLN	A	72	-21.755	78.022	-21.282	1.00	29.17	T3
50	ATOM	2870	N	ARG	A	73	-23.736	77.891	-22.360	1.00	32.66	T3
	ATOM	2871	CA	ARG	A	73	-23.908	79.320	-22.518	1.00	34.99	T3
	ATOM	2872	CB	ARG	A	73	-23.943	79.646	-24.005	1.00	30.30	T3
	ATOM	2873	CG	ARG	A	73	-24.141	81.100	-24.295	1.00	27.65	T3
	ATOM	2874	CD	ARG	A	73	-24.729	81.301	-25.662	1.00	38.36	T3
55	ATOM	2875	NE	ARG	A	73	-24.949	82.713	-25.941	1.00	32.10	T3
	ATOM	2876	CZ	ARG	A	73	-25.712	83.167	-26.929	1.00	37.80	T3
	ATOM	2877	NH1	ARG	A	73	-26.333	82.317	-27.738	1.00	34.69	T3
	ATOM	2878	NH2	ARG	A	73	-25.854	84.472	-27.105	1.00	31.05	T3
	ATOM	2879	C	ARG	A	73	-25.207	79.802	-21.859	1.00	28.70	T3
60	ATOM	2880	O	ARG	A	73	-26.260	79.201	-22.049	1.00	30.68	T3
	ATOM	2881	N	LYS	A	74	-25.125	80.881	-21.082	1.00	30.24	T3
	ATOM	2882	CA	LYS	A	74	-26.303	81.452	-20.432	1.00	28.75	T3
	ATOM	2883	CB	LYS	A	74	-26.028	81.771	-18.967	1.00	32.57	T3
	ATOM	2884	CG	LYS	A	74	-25.814	80.551	-18.085	1.00	29.94	T3
65	ATOM	2885	CD	LYS	A	74	-25.454	80.935	-16.633	1.00	31.92	T3
	ATOM	2886	CE	LYS	A	74	-25.211	79.692	-15.767	1.00	31.71	T3

	ATOM	2887	NZ	LYS	A	74	-24.890	80.021	-14.334	1.00	28.95	T3
	ATOM	2888	C	LYS	A	74	-26.659	82.743	-21.145	1.00	35.63	T3
	ATOM	2889	O	LYS	A	74	-26.038	83.779	-20.905	1.00	34.87	T3
	ATOM	2890	N	LYS	A	75	-27.660	82.677	-22.017	1.00	28.11	T3
5	ATOM	2891	CA	LYS	A	75	-28.100	83.840	-22.780	1.00	28.92	T3
	ATOM	2892	CB	LYS	A	75	-29.258	83.460	-23.707	1.00	27.55	T3
	ATOM	2893	CG	LYS	A	75	-28.961	82.387	-24.737	1.00	38.38	T3
	ATOM	2894	CD	LYS	A	75	-30.202	82.133	-25.582	1.00	26.06	T3
	ATOM	2895	CE	LYS	A	75	-29.951	81.046	-26.627	1.00	30.83	T3
10	ATOM	2896	NZ	LYS	A	75	-31.170	80.668	-27.435	1.00	30.87	T3
	ATOM	2897	C	LYS	A	75	-28.567	84.970	-21.874	1.00	29.15	T3
	ATOM	2898	O	LYS	A	75	-29.287	84.731	-20.912	1.00	22.39	T3
	ATOM	2899	N	VAL	A	76	-28.168	86.199	-22.186	1.00	32.55	T3
	ATOM	2900	CA	VAL	A	76	-28.587	87.352	-21.394	1.00	34.67	T3
15	ATOM	2901	CB	VAL	A	76	-27.764	88.603	-21.681	1.00	29.43	T3
	ATOM	2902	CG1	VAL	A	76	-27.955	89.595	-20.578	1.00	35.39	T3
	ATOM	2903	CG2	VAL	A	76	-26.337	88.256	-21.850	1.00	28.80	T3
	ATOM	2904	C	VAL	A	76	-29.997	87.693	-21.830	1.00	27.72	T3
	ATOM	2905	O	VAL	A	76	-30.852	88.029	-21.015	1.00	33.53	T3
20	ATOM	2906	N	HIS	A	77	-30.216	87.614	-23.135	1.00	28.53	T3
	ATOM	2907	CA	HIS	A	77	-31.501	87.918	-23.721	1.00	29.47	T3
	ATOM	2908	CB	HIS	A	77	-31.309	88.813	-24.938	1.00	29.13	T3
	ATOM	2909	CG	HIS	A	77	-30.610	90.103	-24.631	1.00	25.46	T3
	ATOM	2910	CD2	HIS	A	77	-29.951	90.974	-25.432	1.00	25.67	T3
25	ATOM	2911	ND1	HIS	A	77	-30.604	90.662	-23.370	1.00	32.09	T3
	ATOM	2912	CE1	HIS	A	77	-29.977	91.823	-23.408	1.00	30.94	T3
	ATOM	2913	NE2	HIS	A	77	-29.574	92.035	-24.647	1.00	30.24	T3
	ATOM	2914	C	HIS	A	77	-32.177	86.622	-24.109	1.00	29.26	T3
	ATOM	2915	O	HIS	A	77	-31.503	85.668	-24.503	1.00	32.46	T3
30	ATOM	2916	N	VAL	A	78	-33.508	86.588	-24.017	1.00	34.49	T3
	ATOM	2917	CA	VAL	A	78	-34.213	85.362	-24.317	1.00	25.64	T3
	ATOM	2918	CB	VAL	A	78	-34.755	84.755	-23.011	1.00	30.64	T3
	ATOM	2919	CG1	VAL	A	78	-35.347	83.384	-23.272	1.00	28.67	T3
	ATOM	2920	CG2	VAL	A	78	-33.619	84.607	-22.012	1.00	24.90	T3
35	ATOM	2921	C	VAL	A	78	-35.289	85.328	-25.406	1.00	34.20	T3
	ATOM	2922	O	VAL	A	78	-35.088	84.669	-26.429	1.00	33.52	T3
	ATOM	2923	N	PHE	A	79	-36.417	86.002	-25.227	1.00	27.58	T3
	ATOM	2924	CA	PHE	A	79	-37.481	85.950	-26.260	1.00	27.94	T3
	ATOM	2925	CB	PHE	A	79	-36.920	86.097	-27.689	1.00	27.81	T3
40	ATOM	2926	CG	PHE	A	79	-36.035	87.287	-27.883	1.00	30.87	T3
	ATOM	2927	CD1	PHE	A	79	-34.651	87.146	-27.897	1.00	29.14	T3
	ATOM	2928	CD2	PHE	A	79	-36.582	88.551	-28.047	1.00	29.20	T3
	ATOM	2929	CE1	PHE	A	79	-33.825	88.239	-28.071	1.00	33.33	T3
	ATOM	2930	CE2	PHE	A	79	-35.764	89.659	-28.222	1.00	24.00	T3
45	ATOM	2931	CZ	PHE	A	79	-34.382	89.501	-28.234	1.00	36.57	T3
	ATOM	2932	C	PHE	A	79	-38.326	84.664	-26.269	1.00	30.09	T3
	ATOM	2933	O	PHE	A	79	-37.805	83.559	-26.412	1.00	31.15	T3
	ATOM	2934	N	GLY	A	80	-39.636	84.832	-26.148	1.00	40.66	T3
	ATOM	2935	CA	GLY	A	80	-40.552	83.706	-26.179	1.00	29.41	T3
50	ATOM	2936	C	GLY	A	80	-40.184	82.494	-25.347	1.00	30.86	T3
	ATOM	2937	O	GLY	A	80	-39.842	82.614	-24.168	1.00	30.34	T3
	ATOM	2938	N	ASP	A	81	-40.252	81.320	-25.970	1.00	35.90	T3
	ATOM	2939	CA	ASP	A	81	-39.955	80.067	-25.295	1.00	24.08	T3
	ATOM	2940	CB	ASP	A	81	-40.966	78.994	-25.722	1.00	31.14	T3
55	ATOM	2941	CG	ASP	A	81	-40.847	78.620	-27.182	1.00	28.70	T3
	ATOM	2942	OD1	ASP	A	81	-40.224	79.376	-27.945	1.00	28.39	T3
	ATOM	2943	OD2	ASP	A	81	-41.387	77.569	-27.576	1.00	33.47	T3
	ATOM	2944	C	ASP	A	81	-38.536	79.553	-25.490	1.00	23.40	T3
	ATOM	2945	O	ASP	A	81	-38.299	78.347	-25.464	1.00	25.14	T3
60	ATOM	2946	N	GLU	A	82	-37.585	80.457	-25.686	1.00	32.31	T3
	ATOM	2947	CA	GLU	A	82	-36.196	80.042	-25.844	1.00	32.67	T3
	ATOM	2948	CB	GLU	A	82	-35.303	81.214	-26.223	1.00	27.06	T3
	ATOM	2949	CG	GLU	A	82	-35.308	81.641	-27.651	1.00	27.22	T3
	ATOM	2950	CD	GLU	A	82	-33.987	82.296	-28.016	1.00	32.02	T3
65	ATOM	2951	OE1	GLU	A	82	-33.488	83.112	-27.214	1.00	35.04	T3
	ATOM	2952	OE2	GLU	A	82	-33.439	81.996	-29.097	1.00	32.62	T3

	ATOM	2953	C	GLU	A	82	-35.712	79.568	-24.492	1.00	32.39	T3
	ATOM	2954	O	GLU	A	82	-36.191	80.055	-23.468	1.00	31.91	T3
	ATOM	2955	N	LEU	A	83	-34.773	78.628	-24.472	1.00	23.24	T3
	ATOM	2956	CA	LEU	A	83	-34.211	78.190	-23.199	1.00	30.30	T3
5	ATOM	2957	CB	LEU	A	83	-33.772	76.728	-23.236	1.00	28.43	T3
	ATOM	2958	CG	LEU	A	83	-34.869	75.672	-23.320	1.00	24.55	T3
	ATOM	2959	CD1	LEU	A	83	-35.619	75.822	-24.637	1.00	33.00	T3
	ATOM	2960	CD2	LEU	A	83	-34.253	74.297	-23.220	1.00	29.95	T3
	ATOM	2961	C	LEU	A	83	-32.999	79.085	-23.079	1.00	27.33	T3
10	ATOM	2962	O	LEU	A	83	-32.229	79.209	-24.030	1.00	26.73	T3
	ATOM	2963	N	SER	A	84	-32.834	79.728	-21.931	1.00	34.81	T3
	ATOM	2964	CA	SER	A	84	-31.711	80.633	-21.752	1.00	30.98	T3
	ATOM	2965	CB	SER	A	84	-32.001	81.628	-20.623	1.00	29.30	T3
	ATOM	2966	OG	SER	A	84	-32.490	80.973	-19.473	1.00	30.56	T3
15	ATOM	2967	C	SER	A	84	-30.394	79.922	-21.501	1.00	27.61	T3
	ATOM	2968	O	SER	A	84	-29.348	80.564	-21.417	1.00	29.47	T3
	ATOM	2969	N	LEU	A	85	-30.440	78.598	-21.393	1.00	34.27	T3
	ATOM	2970	CA	LEU	A	85	-29.231	77.821	-21.159	1.00	33.50	T3
	ATOM	2971	CB	LEU	A	85	-29.362	77.023	-19.855	1.00	35.33	T3
20	ATOM	2972	CG	LEU	A	85	-28.119	76.420	-19.173	1.00	31.49	T3
	ATOM	2973	CD1	LEU	A	85	-27.560	75.259	-19.996	1.00	30.74	T3
	ATOM	2974	CD2	LEU	A	85	-27.075	77.502	-18.973	1.00	34.13	T3
	ATOM	2975	C	LEU	A	85	-29.023	76.889	-22.338	1.00	34.19	T3
	ATOM	2976	O	LEU	A	85	-29.748	75.919	-22.501	1.00	29.01	T3
25	ATOM	2977	N	VAL	A	86	-28.044	77.203	-23.176	1.00	31.56	T3
	ATOM	2978	CA	VAL	A	86	-27.741	76.382	-24.344	1.00	29.35	T3
	ATOM	2979	CB	VAL	A	86	-27.577	77.227	-25.618	1.00	32.72	T3
	ATOM	2980	CG1	VAL	A	86	-27.168	76.335	-26.774	1.00	34.52	T3
	ATOM	2981	CG2	VAL	A	86	-28.865	77.957	-25.936	1.00	31.10	T3
30	ATOM	2982	C	VAL	A	86	-26.427	75.681	-24.109	1.00	25.65	T3
	ATOM	2983	O	VAL	A	86	-25.486	76.275	-23.589	1.00	24.58	T3
	ATOM	2984	N	THR	A	87	-26.354	74.415	-24.484	1.00	34.18	T3
	ATOM	2985	CA	THR	A	87	-25.114	73.696	-24.303	1.00	33.85	T3
	ATOM	2986	CB	THR	A	87	-25.333	72.339	-23.579	1.00	33.36	T3
35	ATOM	2987	OG1	THR	A	87	-24.952	71.265	-24.443	1.00	27.09	T3
	ATOM	2988	CG2	THR	A	87	-26.785	72.180	-23.157	1.00	29.85	T3
	ATOM	2989	C	THR	A	87	-24.473	73.498	-25.665	1.00	36.00	T3
	ATOM	2990	O	THR	A	87	-25.049	72.886	-26.559	1.00	32.74	T3
	ATOM	2991	N	LEU	A	88	-23.285	74.068	-25.820	1.00	32.26	T3
40	ATOM	2992	CA	LEU	A	88	-22.514	73.985	-27.057	1.00	30.19	T3
	ATOM	2993	CB	LEU	A	88	-21.743	75.292	-27.275	1.00	36.43	T3
	ATOM	2994	CG	LEU	A	88	-22.372	76.690	-27.319	1.00	39.53	T3
	ATOM	2995	CD1	LEU	A	88	-23.614	76.763	-26.501	1.00	39.12	T3
	ATOM	2996	CD2	LEU	A	88	-21.378	77.685	-26.793	1.00	30.61	T3
45	ATOM	2997	C	LEU	A	88	-21.491	72.865	-26.873	1.00	28.47	T3
	ATOM	2998	O	LEU	A	88	-21.008	72.627	-25.763	1.00	30.43	T3
	ATOM	2999	N	PHE	A	89	-21.173	72.140	-27.930	1.00	29.92	T3
	ATOM	3000	CA	PHE	A	89	-20.120	71.123	-27.811	1.00	36.27	T3
	ATOM	3001	CB	PHE	A	89	-18.828	71.824	-27.378	1.00	31.51	T3
50	ATOM	3002	CG	PHE	A	89	-18.603	73.120	-28.094	1.00	35.86	T3
	ATOM	3003	CD1	PHE	A	89	-18.008	74.200	-27.453	1.00	38.91	T3
	ATOM	3004	CD2	PHE	A	89	-19.087	73.290	-29.406	1.00	26.49	T3
	ATOM	3005	CE1	PHE	A	89	-17.911	75.440	-28.104	1.00	33.28	T3
	ATOM	3006	CE2	PHE	A	89	-18.995	74.522	-30.064	1.00	34.72	T3
55	ATOM	3007	CZ	PHE	A	89	-18.409	75.601	-29.412	1.00	39.34	T3
	ATOM	3008	C	PHE	A	89	-20.356	69.886	-26.943	1.00	25.51	T3
	ATOM	3009	O	PHE	A	89	-21.065	68.983	-27.372	1.00	29.29	T3
	ATOM	3010	N	ARG	A	90	-19.751	69.805	-25.756	1.00	28.37	T3
	ATOM	3011	CA	ARG	A	90	-19.939	68.600	-24.928	1.00	24.36	T3
60	ATOM	3012	CB	ARG	A	90	-21.437	68.327	-24.729	1.00	29.88	T3
	ATOM	3013	CG	ARG	A	90	-21.784	66.860	-24.518	1.00	26.11	T3
	ATOM	3014	CD	ARG	A	90	-23.264	66.615	-24.719	1.00	25.22	T3
	ATOM	3015	NE	ARG	A	90	-23.542	65.183	-24.730	1.00	28.21	T3
	ATOM	3016	CZ	ARG	A	90	-24.443	64.598	-25.518	1.00	26.76	T3
65	ATOM	3017	NH1	ARG	A	90	-25.172	65.325	-26.367	1.00	23.47	T3
	ATOM	3018	NH2	ARG	A	90	-24.590	63.277	-25.478	1.00	26.31	T3

	ATOM	3019	C	ARG	A	90	-19.293	67.320	-25.508	1.00	23.23	T3
	ATOM	3020	O	ARG	A	90	-19.540	66.957	-26.664	1.00	30.57	T3
	ATOM	3021	N	CYS	A	91	-18.499	66.623	-24.690	1.00	29.07	T3
	ATOM	3022	CA	CYS	A	91	-17.836	65.383	-25.119	1.00	34.20	T3
5	ATOM	3023	CB	CYS	A	91	-16.343	65.626	-25.289	1.00	29.93	T3
	ATOM	3024	SG	CYS	A	91	-15.593	66.287	-23.798	1.00	26.71	T3
	ATOM	3025	C	CYS	A	91	-18.061	64.227	-24.133	1.00	35.24	T3
	ATOM	3026	O	CYS	A	91	-18.521	64.436	-23.009	1.00	28.19	T3
	ATOM	3027	N	ILE	A	92	-17.727	63.011	-24.559	1.00	31.76	T3
10	ATOM	3028	CA	ILE	A	92	-17.923	61.815	-23.734	1.00	27.31	T3
	ATOM	3029	CB	ILE	A	92	-19.189	61.039	-24.172	1.00	30.68	T3
	ATOM	3030	CG2	ILE	A	92	-19.403	59.840	-23.277	1.00	32.97	T3
	ATOM	3031	CG1	ILE	A	92	-20.419	61.952	-24.115	1.00	33.61	T3
	ATOM	3032	CD1	ILE	A	92	-21.637	61.381	-24.837	1.00	29.23	T3
15	ATOM	3033	C	ILE	A	92	-16.744	60.868	-23.895	1.00	31.58	T3
	ATOM	3034	O	ILE	A	92	-16.077	60.884	-24.917	1.00	29.95	T3
	ATOM	3035	N	GLN	A	93	-16.493	60.039	-22.889	1.00	29.53	T3
	ATOM	3036	CA	GLN	A	93	-15.401	59.068	-22.947	1.00	33.04	T3
	ATOM	3037	CB	GLN	A	93	-14.095	59.691	-22.455	1.00	39.34	T3
20	ATOM	3038	CG	GLN	A	93	-13.316	60.428	-23.514	1.00	37.37	T3
	ATOM	3039	CD	GLN	A	93	-12.211	59.585	-24.118	1.00	33.41	T3
	ATOM	3040	OE1	GLN	A	93	-12.455	58.680	-24.919	1.00	30.61	T3
	ATOM	3041	NE2	GLN	A	93	-10.977	59.877	-23.725	1.00	36.52	T3
	ATOM	3042	C	GLN	A	93	-15.718	57.836	-22.104	1.00	25.66	T3
25	ATOM	3043	O	GLN	A	93	-15.958	57.940	-20.886	1.00	24.53	T3
	ATOM	3044	N	ASN	A	94	-15.746	56.672	-22.747	1.00	32.35	T3
	ATOM	3045	CA	ASN	A	94	-16.008	55.443	-22.014	1.00	27.93	T3
	ATOM	3046	CB	ASN	A	94	-16.027	54.238	-22.965	1.00	26.64	T3
	ATOM	3047	CG	ASN	A	94	-17.347	54.090	-23.692	1.00	37.32	T3
30	ATOM	3048	OD1	ASN	A	94	-18.400	54.034	-23.067	1.00	36.65	T3
	ATOM	3049	ND2	ASN	A	94	-17.296	54.014	-25.015	1.00	25.72	T3
	ATOM	3050	C	ASN	A	94	-14.866	55.303	-20.999	1.00	30.66	T3
	ATOM	3051	O	ASN	A	94	-13.726	55.712	-21.272	1.00	32.20	T3
	ATOM	3052	N	MET	A	95	-15.172	54.751	-19.830	1.00	38.37	T3
35	ATOM	3053	CA	MET	A	95	-14.163	54.572	-18.797	1.00	31.38	T3
	ATOM	3054	CB	MET	A	95	-14.610	55.231	-17.486	1.00	28.77	T3
	ATOM	3055	CG	MET	A	95	-14.823	56.727	-17.584	1.00	28.46	T3
	ATOM	3056	SD	MET	A	95	-13.400	57.581	-18.306	1.00	33.28	T3
	ATOM	3057	CE	MET	A	95	-12.210	57.555	-16.938	1.00	26.16	T3
40	ATOM	3058	C	MET	A	95	-13.917	53.091	-18.544	1.00	22.16	T3
	ATOM	3059	O	MET	A	95	-14.813	52.261	-18.748	1.00	33.34	T3
	ATOM	3060	N	PRO	A	96	-12.689	52.739	-18.110	1.00	27.33	T3
	ATOM	3061	CD	PRO	A	96	-11.513	53.613	-17.991	1.00	23.44	T3
	ATOM	3062	CA	PRO	A	96	-12.313	51.355	-17.814	1.00	34.68	T3
45	ATOM	3063	CB	PRO	A	96	-10.784	51.396	-17.743	1.00	28.02	T3
	ATOM	3064	CG	PRO	A	96	-10.412	52.671	-18.411	1.00	34.40	T3
	ATOM	3065	C	PRO	A	96	-12.897	51.074	-16.445	1.00	22.41	T3
	ATOM	3066	O	PRO	A	96	-13.664	51.888	-15.902	1.00	35.85	T3
	ATOM	3067	N	GLU	A	97	-12.514	49.947	-15.861	1.00	31.14	T3
50	ATOM	3068	CA	GLU	A	97	-13.028	49.605	-14.547	1.00	32.77	T3
	ATOM	3069	CB	GLU	A	97	-13.671	48.229	-14.602	1.00	33.95	T3
	ATOM	3070	CG	GLU	A	97	-14.551	47.950	-13.417	1.00	35.98	T3
	ATOM	3071	CD	GLU	A	97	-15.858	47.314	-13.840	1.00	25.83	T3
	ATOM	3072	OE1	GLU	A	97	-15.809	46.210	-14.451	1.00	32.30	T3
55	ATOM	3073	OE2	GLU	A	97	-16.935	47.920	-13.574	1.00	31.72	T3
	ATOM	3074	C	GLU	A	97	-11.907	49.615	-13.522	1.00	27.24	T3
	ATOM	3075	O	GLU	A	97	-12.137	49.791	-12.321	1.00	32.06	T3
	ATOM	3076	N	THR	A	98	-10.688	49.458	-14.019	1.00	27.47	T3
	ATOM	3077	CA	THR	A	98	-9.522	49.400	-13.158	1.00	40.96	T3
60	ATOM	3078	CB	THR	A	98	-8.507	48.406	-13.711	1.00	36.56	T3
	ATOM	3079	OG1	THR	A	98	-8.047	48.877	-14.988	1.00	26.86	T3
	ATOM	3080	CG2	THR	A	98	-9.155	47.026	-13.876	1.00	24.00	T3
	ATOM	3081	C	THR	A	98	-8.799	50.717	-12.909	1.00	32.97	T3
	ATOM	3082	O	THR	A	98	-8.880	51.268	-11.810	1.00	37.02	T3
65	ATOM	3083	N	LEU	A	99	-8.091	51.232	-13.908	1.00	28.16	T3
	ATOM	3084	CA	LEU	A	99	-7.350	52.461	-13.681	1.00	37.74	T3

	ATOM	3085	CB	LEU	A	99	-5.867	52.193	-13.893	1.00	36.74	T3
	ATOM	3086	CG	LEU	A	99	-5.330	51.161	-12.899	1.00	27.88	T3
	ATOM	3087	CD1	LEU	A	99	-3.891	50.783	-13.229	1.00	24.33	T3
	ATOM	3088	CD2	LEU	A	99	-5.419	51.737	-11.504	1.00	28.15	T3
5	ATOM	3089	C	LEU	A	99	-7.801	53.650	-14.514	1.00	31.40	T3
	ATOM	3090	O	LEU	A	99	-7.096	54.087	-15.430	1.00	26.10	T3
	ATOM	3091	N	PRO	A	100	-8.977	54.211	-14.187	1.00	32.19	T3
	ATOM	3092	CD	PRO	A	100	-9.823	53.842	-13.039	1.00	32.97	T3
	ATOM	3093	CA	PRO	A	100	-9.553	55.357	-14.889	1.00	27.14	T3
10	ATOM	3094	CB	PRO	A	100	-10.669	55.804	-13.943	1.00	27.00	T3
	ATOM	3095	CG	PRO	A	100	-11.142	54.516	-13.384	1.00	29.64	T3
	ATOM	3096	C	PRO	A	100	-8.543	56.461	-15.155	1.00	32.39	T3
	ATOM	3097	O	PRO	A	100	-7.916	56.992	-14.237	1.00	29.94	T3
	ATOM	3098	N	ASN	A	101	-8.402	56.801	-16.426	1.00	35.51	T3
15	ATOM	3099	CA	ASN	A	101	-7.477	57.844	-16.848	1.00	20.74	T3
	ATOM	3100	CB	ASN	A	101	-6.039	57.338	-16.813	1.00	32.86	T3
	ATOM	3101	CG	ASN	A	101	-5.411	57.513	-15.471	1.00	33.61	T3
	ATOM	3102	OD1	ASN	A	101	-5.160	58.645	-15.041	1.00	32.71	T3
	ATOM	3103	ND2	ASN	A	101	-5.164	56.400	-14.778	1.00	36.16	T3
20	ATOM	3104	C	ASN	A	101	-7.781	58.255	-18.262	1.00	29.77	T3
	ATOM	3105	O	ASN	A	101	-7.115	57.778	-19.192	1.00	34.42	T3
	ATOM	3106	N	ASN	A	102	-8.762	59.133	-18.457	1.00	33.77	T3
	ATOM	3107	CA	ASN	A	102	-9.035	59.491	-19.829	1.00	29.61	T3
	ATOM	3108	CB	ASN	A	102	-10.400	58.950	-20.245	1.00	30.67	T3
25	ATOM	3109	CG	ASN	A	102	-10.314	57.485	-20.689	1.00	25.19	T3
	ATOM	3110	OD1	ASN	A	102	-9.411	57.105	-21.449	1.00	37.29	T3
	ATOM	3111	ND2	ASN	A	102	-11.247	56.660	-20.213	1.00	32.84	T3
	ATOM	3112	C	ASN	A	102	-8.814	60.901	-20.357	1.00	30.40	T3
	ATOM	3113	O	ASN	A	102	-8.190	61.042	-21.417	1.00	30.57	T3
30	ATOM	3114	N	SER	A	103	-9.267	61.949	-19.681	1.00	30.89	T3
	ATOM	3115	CA	SER	A	103	-9.006	63.270	-20.265	1.00	33.14	T3
	ATOM	3116	CB	SER	A	103	-7.485	63.501	-20.388	1.00	28.67	T3
	ATOM	3117	OG	SER	A	103	-7.110	63.979	-21.678	1.00	30.98	T3
	ATOM	3118	C	SER	A	103	-9.665	63.406	-21.653	1.00	26.65	T3
35	ATOM	3119	O	SER	A	103	-9.510	62.563	-22.543	1.00	29.20	T3
	ATOM	3120	N	CYS	A	104	-10.395	64.488	-21.846	1.00	28.36	T3
	ATOM	3121	CA	CYS	A	104	-11.061	64.680	-23.110	1.00	27.21	T3
	ATOM	3122	CB	CYS	A	104	-12.457	64.081	-23.018	1.00	32.83	T3
	ATOM	3123	SG	CYS	A	104	-13.337	63.975	-24.557	1.00	30.81	T3
40	ATOM	3124	C	CYS	A	104	-11.125	66.165	-23.432	1.00	36.61	T3
	ATOM	3125	O	CYS	A	104	-11.565	66.977	-22.618	1.00	38.21	T3
	ATOM	3126	N	TYR	A	105	-10.665	66.510	-24.626	1.00	35.65	T3
	ATOM	3127	CA	TYR	A	105	-10.657	67.889	-25.086	1.00	32.23	T3
	ATOM	3128	CB	TYR	A	105	-9.291	68.220	-25.697	1.00	28.49	T3
45	ATOM	3129	CG	TYR	A	105	-9.187	69.600	-26.311	1.00	34.58	T3
	ATOM	3130	CD1	TYR	A	105	-8.697	70.674	-25.574	1.00	28.28	T3
	ATOM	3131	CE1	TYR	A	105	-8.577	71.939	-26.141	1.00	35.78	T3
	ATOM	3132	CD2	TYR	A	105	-9.563	69.826	-27.635	1.00	28.05	T3
	ATOM	3133	CE2	TYR	A	105	-9.450	71.088	-28.208	1.00	29.64	T3
50	ATOM	3134	CZ	TYR	A	105	-8.954	72.137	-27.458	1.00	33.13	T3
	ATOM	3135	OH	TYR	A	105	-8.819	73.377	-28.030	1.00	27.94	T3
	ATOM	3136	C	TYR	A	105	-11.733	68.077	-26.147	1.00	27.39	T3
	ATOM	3137	O	TYR	A	105	-12.027	67.169	-26.919	1.00	32.23	T3
	ATOM	3138	N	SER	A	106	-12.318	69.261	-26.182	1.00	33.49	T3
55	ATOM	3139	CA	SER	A	106	-13.327	69.563	-27.178	1.00	29.03	T3
	ATOM	3140	CB	SER	A	106	-14.676	68.968	-26.773	1.00	29.58	T3
	ATOM	3141	OG	SER	A	106	-15.611	69.042	-27.836	1.00	32.60	T3
	ATOM	3142	C	SER	A	106	-13.403	71.077	-27.267	1.00	28.89	T3
	ATOM	3143	O	SER	A	106	-13.239	71.771	-26.262	1.00	28.26	T3
60	ATOM	3144	N	ALA	A	107	-13.628	71.583	-28.474	1.00	32.23	T3
	ATOM	3145	CA	ALA	A	107	-13.714	73.021	-28.691	1.00	29.90	T3
	ATOM	3146	CB	ALA	A	107	-12.320	73.603	-28.842	1.00	26.56	T3
	ATOM	3147	C	ALA	A	107	-14.550	73.350	-29.920	1.00	29.79	T3
	ATOM	3148	O	ALA	A	107	-14.824	72.486	-30.738	1.00	28.19	T3
65	ATOM	3149	N	GLY	A	108	-14.951	74.609	-30.039	1.00	25.02	T3
	ATOM	3150	CA	GLY	A	108	-15.742	75.034	-31.181	1.00	27.40	T3

	ATOM	3151	C	GLY A 108	-15.924	76.539	-31.203	1.00	26.92	T3
	ATOM	3152	O	GLY A 108	-15.432	77.231	-30.316	1.00	34.23	T3
	ATOM	3153	N	ILE A 109	-16.626	77.049	-32.207	1.00	35.25	T3
	ATOM	3154	CA	ILE A 109	-16.864	78.479	-32.321	1.00	32.07	T3
5	ATOM	3155	CB	ILE A 109	-16.478	79.000	-33.706	1.00	34.44	T3
	ATOM	3156	CG2	ILE A 109	-16.773	80.481	-33.797	1.00	24.01	T3
	ATOM	3157	CG1	ILE A 109	-14.998	78.735	-33.968	1.00	29.21	T3
	ATOM	3158	CD1	ILE A 109	-14.581	79.001	-35.383	1.00	33.14	T3
	ATOM	3159	C	ILE A 109	-18.336	78.765	-32.104	1.00	32.48	T3
10	ATOM	3160	O	ILE A 109	-19.188	78.006	-32.548	1.00	27.08	T3
	ATOM	3161	N	ALA A 110	-18.635	79.861	-31.418	1.00	22.56	T3
	ATOM	3162	CA	ALA A 110	-20.016	80.237	-31.148	1.00	33.40	T3
	ATOM	3163	CB	ALA A 110	-20.473	79.634	-29.845	1.00	23.85	T3
	ATOM	3164	C	ALA A 110	-20.128	81.743	-31.075	1.00	29.17	T3
15	ATOM	3165	O	ALA A 110	-19.151	82.426	-30.779	1.00	28.07	T3
	ATOM	3166	N	LYS A 111	-21.313	82.270	-31.359	1.00	29.80	T3
	ATOM	3167	CA	LYS A 111	-21.499	83.708	-31.278	1.00	27.23	T3
	ATOM	3168	CB	LYS A 111	-22.455	84.207	-32.350	1.00	31.18	T3
	ATOM	3169	CG	LYS A 111	-22.603	85.721	-32.314	1.00	28.64	T3
20	ATOM	3170	CD	LYS A 111	-23.521	86.231	-33.417	1.00	32.17	T3
	ATOM	3171	CE	LYS A 111	-23.646	87.763	-33.387	1.00	33.16	T3
	ATOM	3172	NZ	LYS A 111	-24.520	88.292	-34.496	1.00	34.59	T3
	ATOM	3173	C	LYS A 111	-22.066	84.009	-29.901	1.00	25.32	T3
	ATOM	3174	O	LYS A 111	-23.000	83.346	-29.472	1.00	27.20	T3
25	ATOM	3175	N	LEU A 112	-21.489	84.994	-29.215	1.00	28.88	T3
	ATOM	3176	CA	LEU A 112	-21.914	85.373	-27.872	1.00	32.04	T3
	ATOM	3177	CB	LEU A 112	-20.822	85.008	-26.868	1.00	36.54	T3
	ATOM	3178	CG	LEU A 112	-20.249	83.598	-26.952	1.00	19.81	T3
	ATOM	3179	CD1	LEU A 112	-18.985	83.502	-26.120	1.00	28.10	T3
30	ATOM	3180	CD2	LEU A 112	-21.288	82.614	-26.488	1.00	31.91	T3
	ATOM	3181	C	LEU A 112	-22.183	86.872	-27.781	1.00	33.42	T3
	ATOM	3182	O	LEU A 112	-21.718	87.641	-28.624	1.00	30.72	T3
	ATOM	3183	N	GLU A 113	-22.917	87.286	-26.749	1.00	26.61	T3
	ATOM	3184	CA	GLU A 113	-23.223	88.696	-26.551	1.00	35.48	T3
35	ATOM	3185	CB	GLU A 113	-24.701	88.973	-26.743	1.00	35.36	T3
	ATOM	3186	CG	GLU A 113	-25.378	88.236	-27.848	1.00	28.78	T3
	ATOM	3187	CD	GLU A 113	-26.692	88.913	-28.239	1.00	29.57	T3
	ATOM	3188	OE1	GLU A 113	-27.475	89.315	-27.322	1.00	31.35	T3
	ATOM	3189	OE2	GLU A 113	-26.939	89.040	-29.472	1.00	32.71	T3
40	ATOM	3190	C	GLU A 113	-22.881	89.184	-25.159	1.00	30.67	T3
	ATOM	3191	O	GLU A 113	-22.801	88.403	-24.220	1.00	31.57	T3
	ATOM	3192	N	GLU A 114	-22.708	90.495	-25.032	1.00	31.32	T3
	ATOM	3193	CA	GLU A 114	-22.417	91.113	-23.746	1.00	30.76	T3
	ATOM	3194	CB	GLU A 114	-22.716	92.601	-23.782	1.00	30.58	T3
45	ATOM	3195	CG	GLU A 114	-21.614	93.494	-24.211	1.00	28.49	T3
	ATOM	3196	CD	GLU A 114	-21.885	94.916	-23.762	1.00	27.42	T3
	ATOM	3197	OE1	GLU A 114	-21.930	95.147	-22.528	1.00	36.04	T3
	ATOM	3198	OE2	GLU A 114	-22.065	95.794	-24.637	1.00	29.25	T3
	ATOM	3199	C	GLU A 114	-23.332	90.531	-22.689	1.00	31.97	T3
50	ATOM	3200	O	GLU A 114	-24.548	90.566	-22.839	1.00	31.37	T3
	ATOM	3201	N	GLY A 115	-22.766	90.020	-21.609	1.00	27.87	T3
	ATOM	3202	CA	GLY A 115	-23.602	89.492	-20.555	1.00	30.27	T3
	ATOM	3203	C	GLY A 115	-23.741	87.993	-20.567	1.00	32.51	T3
	ATOM	3204	O	GLY A 115	-24.147	87.413	-19.563	1.00	31.37	T3
55	ATOM	3205	N	ASP A 116	-23.440	87.350	-21.692	1.00	35.92	T3
	ATOM	3206	CA	ASP A 116	-23.541	85.900	-21.731	1.00	33.81	T3
	ATOM	3207	CB	ASP A 116	-23.300	85.355	-23.144	1.00	28.70	T3
	ATOM	3208	CG	ASP A 116	-24.434	85.660	-24.106	1.00	31.92	T3
	ATOM	3209	OD1	ASP A 116	-25.582	85.826	-23.642	1.00	29.93	T3
60	ATOM	3210	OD2	ASP A 116	-24.179	85.711	-25.333	1.00	31.86	T3
	ATOM	3211	C	ASP A 116	-22.467	85.350	-20.797	1.00	24.30	T3
	ATOM	3212	O	ASP A 116	-21.441	85.997	-20.556	1.00	34.44	T3
	ATOM	3213	N	GLU A 117	-22.704	84.164	-20.256	1.00	31.55	T3
	ATOM	3214	CA	GLU A 117	-21.724	83.546	-19.382	1.00	25.60	T3
65	ATOM	3215	CB	GLU A 117	-22.239	83.478	-17.953	1.00	27.61	T3
	ATOM	3216	CG	GLU A 117	-22.687	84.798	-17.397	1.00	31.94	T3

	ATOM	3217	CD	GLU	A	117	-23.112	84.688	-15.939	1.00	29.30	T3
	ATOM	3218	OE1	GLU	A	117	-23.722	83.651	-15.568	1.00	30.47	T3
	ATOM	3219	OE2	GLU	A	117	-22.847	85.645	-15.167	1.00	28.83	T3
	ATOM	3220	C	GLU	A	117	-21.455	82.136	-19.893	1.00	29.57	T3
5	ATOM	3221	O	GLU	A	117	-22.369	81.446	-20.355	1.00	30.63	T3
	ATOM	3222	N	LEU	A	118	-20.193	81.726	-19.830	1.00	26.16	T3
	ATOM	3223	CA	LEU	A	118	-19.801	80.395	-20.262	1.00	23.68	T3
	ATOM	3224	CB	LEU	A	118	-18.596	80.460	-21.193	1.00	37.23	T3
	ATOM	3225	CG	LEU	A	118	-18.795	81.145	-22.534	1.00	24.34	T3
10	ATOM	3226	CD1	LEU	A	118	-17.525	81.050	-23.338	1.00	36.55	T3
	ATOM	3227	CD2	LEU	A	118	-19.934	80.480	-23.270	1.00	33.02	T3
	ATOM	3228	C	LEU	A	118	-19.427	79.601	-19.025	1.00	28.91	T3
	ATOM	3229	O	LEU	A	118	-18.892	80.156	-18.063	1.00	33.31	T3
	ATOM	3230	N	GLN	A	119	-19.714	78.304	-19.047	1.00	26.79	T3
15	ATOM	3231	CA	GLN	A	119	-19.376	77.438	-17.926	1.00	37.34	T3
	ATOM	3232	CB	GLN	A	119	-20.491	77.470	-16.897	1.00	30.54	T3
	ATOM	3233	CG	GLN	A	119	-21.801	76.935	-17.409	1.00	30.53	T3
	ATOM	3234	CD	GLN	A	119	-22.916	77.066	-16.387	1.00	22.96	T3
	ATOM	3235	OE1	GLN	A	119	-23.957	76.408	-16.493	1.00	28.56	T3
20	ATOM	3236	NE2	GLN	A	119	-22.713	77.927	-15.397	1.00	24.99	T3
	ATOM	3237	C	GLN	A	119	-19.122	76.003	-18.380	1.00	28.93	T3
	ATOM	3238	O	GLN	A	119	-19.637	75.565	-19.408	1.00	26.69	T3
	ATOM	3239	N	LEU	A	120	-18.316	75.287	-17.604	1.00	32.43	T3
	ATOM	3240	CA	LEU	A	120	-17.946	73.900	-17.885	1.00	32.14	T3
25	ATOM	3241	CB	LEU	A	120	-16.426	73.751	-17.750	1.00	30.01	T3
	ATOM	3242	CG	LEU	A	120	-15.623	72.541	-18.217	1.00	34.76	T3
	ATOM	3243	CD1	LEU	A	120	-16.281	71.264	-17.791	1.00	23.57	T3
	ATOM	3244	CD2	LEU	A	120	-15.485	72.590	-19.712	1.00	31.45	T3
	ATOM	3245	C	LEU	A	120	-18.646	73.030	-16.843	1.00	31.62	T3
30	ATOM	3246	O	LEU	A	120	-18.424	73.197	-15.639	1.00	32.09	T3
	ATOM	3247	N	ALA	A	121	-19.481	72.097	-17.285	1.00	38.31	T3
	ATOM	3248	CA	ALA	A	121	-20.196	71.254	-16.339	1.00	34.91	T3
	ATOM	3249	CB	ALA	A	121	-21.660	71.662	-16.316	1.00	34.11	T3
	ATOM	3250	C	ALA	A	121	-20.084	69.747	-16.566	1.00	29.76	T3
35	ATOM	3251	O	ALA	A	121	-20.094	69.265	-17.699	1.00	28.10	T3
	ATOM	3252	N	ILE	A	122	-19.976	69.007	-15.468	1.00	26.47	T3
	ATOM	3253	CA	ILE	A	122	-19.915	67.552	-15.525	1.00	37.40	T3
	ATOM	3254	CB	ILE	A	122	-18.780	67.003	-14.663	1.00	28.90	T3
	ATOM	3255	CG2	ILE	A	122	-18.715	65.493	-14.808	1.00	23.08	T3
40	ATOM	3256	CG1	ILE	A	122	-17.459	67.628	-15.103	1.00	29.70	T3
	ATOM	3257	CD1	ILE	A	122	-16.267	67.152	-14.318	1.00	28.96	T3
	ATOM	3258	C	ILE	A	122	-21.246	66.986	-15.012	1.00	38.11	T3
	ATOM	3259	O	ILE	A	122	-21.622	67.189	-13.855	1.00	27.78	T3
	ATOM	3260	N	PRO	A	123	-21.978	66.270	-15.880	1.00	33.65	T3
45	ATOM	3261	CD	PRO	A	123	-21.654	66.073	-17.301	1.00	35.46	T3
	ATOM	3262	CA	PRO	A	123	-23.276	65.663	-15.560	1.00	32.06	T3
	ATOM	3263	CB	PRO	A	123	-23.821	65.251	-16.933	1.00	20.49	T3
	ATOM	3264	CG	PRO	A	123	-23.022	66.077	-17.911	1.00	37.95	T3
	ATOM	3265	C	PRO	A	123	-23.183	64.460	-14.611	1.00	31.00	T3
50	ATOM	3266	O	PRO	A	123	-23.649	63.364	-14.935	1.00	29.71	T3
	ATOM	3267	N	ARG	A	124	-22.567	64.662	-13.453	1.00	34.09	T3
	ATOM	3268	CA	ARG	A	124	-22.442	63.600	-12.464	1.00	35.13	T3
	ATOM	3269	CB	ARG	A	124	-21.158	62.817	-12.652	1.00	34.60	T3
	ATOM	3270	CG	ARG	A	124	-21.292	61.698	-13.652	1.00	27.46	T3
55	ATOM	3271	CD	ARG	A	124	-20.613	60.457	-13.101	1.00	38.15	T3
	ATOM	3272	NE	ARG	A	124	-21.369	59.845	-12.006	1.00	40.90	T3
	ATOM	3273	CZ	ARG	A	124	-20.897	58.861	-11.241	1.00	29.75	T3
	ATOM	3274	NH1	ARG	A	124	-19.668	58.394	-11.456	1.00	32.36	T3
	ATOM	3275	NH2	ARG	A	124	-21.651	58.332	-10.273	1.00	39.93	T3
60	ATOM	3276	C	ARG	A	124	-22.449	64.210	-11.091	1.00	31.55	T3
	ATOM	3277	O	ARG	A	124	-22.116	65.387	-10.936	1.00	33.01	T3
	ATOM	3278	N	GLU	A	125	-22.813	63.415	-10.091	1.00	30.00	T3
	ATOM	3279	CA	GLU	A	125	-22.885	63.941	-8.742	1.00	30.74	T3
	ATOM	3280	CB	GLU	A	125	-23.734	63.029	-7.870	1.00	31.06	T3
65	ATOM	3281	CG	GLU	A	125	-25.217	63.130	-8.217	1.00	29.61	T3
	ATOM	3282	CD	GLU	A	125	-26.107	63.153	-6.970	1.00	26.80	T3

	ATOM	3283	OE1	GLU	A	125	-26.005	62.201	-6.146	1.00	27.72	T3
	ATOM	3284	OE2	GLU	A	125	-26.905	64.120	-6.816	1.00	29.99	T3
	ATOM	3285	C	GLU	A	125	-21.545	64.227	-8.085	1.00	33.93	T3
	ATOM	3286	O	GLU	A	125	-21.342	65.323	-7.552	1.00	35.98	T3
5	ATOM	3287	N	ASN	A	126	-20.631	63.267	-8.096	1.00	31.11	T3
	ATOM	3288	CA	ASN	A	126	-19.325	63.535	-7.506	1.00	32.66	T3
	ATOM	3289	CB	ASN	A	126	-19.204	62.933	-6.112	1.00	21.97	T3
	ATOM	3290	CG	ASN	A	126	-19.850	63.810	-5.045	1.00	34.56	T3
	ATOM	3291	OD1	ASN	A	126	-21.075	63.811	-4.880	1.00	34.54	T3
10	ATOM	3292	ND2	ASN	A	126	-19.026	64.577	-4.323	1.00	29.03	T3
	ATOM	3293	C	ASN	A	126	-18.268	62.974	-8.412	1.00	36.93	T3
	ATOM	3294	O	ASN	A	126	-17.524	62.057	-8.046	1.00	30.26	T3
	ATOM	3295	N	ALA	A	127	-18.221	63.539	-9.613	1.00	24.06	T3
	ATOM	3296	CA	ALA	A	127	-17.277	63.110	-10.633	1.00	28.58	T3
15	ATOM	3297	CB	ALA	A	127	-17.298	64.094	-11.805	1.00	21.63	T3
	ATOM	3298	C	ALA	A	127	-15.867	62.978	-10.088	1.00	33.21	T3
	ATOM	3299	O	ALA	A	127	-15.378	63.866	-9.391	1.00	25.59	T3
	ATOM	3300	N	GLN	A	128	-15.230	61.850	-10.388	1.00	33.27	T3
	ATOM	3301	CA	GLN	A	128	-13.853	61.626	-9.966	1.00	29.84	T3
20	ATOM	3302	CB	GLN	A	128	-13.568	60.130	-9.870	1.00	37.12	T3
	ATOM	3303	CG	GLN	A	128	-14.360	59.458	-8.760	1.00	35.24	T3
	ATOM	3304	CD	GLN	A	128	-14.303	60.252	-7.455	1.00	33.54	T3
	ATOM	3305	OE1	GLN	A	128	-13.223	60.499	-6.904	1.00	24.75	T3
	ATOM	3306	NE2	GLN	A	128	-15.468	60.663	-6.961	1.00	37.96	T3
25	ATOM	3307	C	GLN	A	128	-12.988	62.290	-11.036	1.00	29.17	T3
	ATOM	3308	O	GLN	A	128	-12.751	61.733	-12.122	1.00	25.78	T3
	ATOM	3309	N	ILE	A	129	-12.520	63.493	-10.707	1.00	26.37	T3
	ATOM	3310	CA	ILE	A	129	-11.748	64.323	-11.627	1.00	26.98	T3
	ATOM	3311	CB	ILE	A	129	-12.585	65.623	-11.908	1.00	28.96	T3
30	ATOM	3312	CG2	ILE	A	129	-11.773	66.874	-11.703	1.00	33.76	T3
	ATOM	3313	CG1	ILE	A	129	-13.206	65.535	-13.292	1.00	31.09	T3
	ATOM	3314	CD1	ILE	A	129	-14.130	64.348	-13.449	1.00	35.22	T3
	ATOM	3315	C	ILE	A	129	-10.353	64.676	-11.125	1.00	28.48	T3
	ATOM	3316	O	ILE	A	129	-10.080	64.606	-9.930	1.00	28.17	T3
35	ATOM	3317	N	SER	A	130	-9.464	65.032	-12.045	1.00	29.34	T3
	ATOM	3318	CA	SER	A	130	-8.120	65.449	-11.661	1.00	32.58	T3
	ATOM	3319	CB	SER	A	130	-7.079	64.890	-12.617	1.00	30.12	T3
	ATOM	3320	OG	SER	A	130	-5.830	65.512	-12.380	1.00	33.17	T3
	ATOM	3321	C	SER	A	130	-8.094	66.973	-11.734	1.00	26.61	T3
40	ATOM	3322	O	SER	A	130	-8.430	67.545	-12.770	1.00	34.58	T3
	ATOM	3323	N	LEU	A	131	-7.701	67.635	-10.649	1.00	36.20	T3
	ATOM	3324	CA	LEU	A	131	-7.663	69.094	-10.651	1.00	24.80	T3
	ATOM	3325	CB	LEU	A	131	-8.185	69.644	-9.321	1.00	23.74	T3
	ATOM	3326	CG	LEU	A	131	-9.704	69.666	-9.173	1.00	32.00	T3
45	ATOM	3327	CD1	LEU	A	131	-10.260	68.269	-9.298	1.00	29.77	T3
	ATOM	3328	CD2	LEU	A	131	-10.060	70.259	-7.840	1.00	26.77	T3
	ATOM	3329	C	LEU	A	131	-6.296	69.700	-10.946	1.00	34.79	T3
	ATOM	3330	O	LEU	A	131	-5.958	70.768	-10.430	1.00	27.38	T3
	ATOM	3331	N	ASP	A	132	-5.511	69.026	-11.778	1.00	30.92	T3
50	ATOM	3332	CA	ASP	A	132	-4.192	69.531	-12.133	1.00	30.74	T3
	ATOM	3333	CB	ASP	A	132	-3.257	68.372	-12.484	1.00	36.98	T3
	ATOM	3334	CG	ASP	A	132	-2.633	67.726	-11.245	1.00	27.18	T3
	ATOM	3335	OD1	ASP	A	132	-2.033	66.629	-11.377	1.00	27.26	T3
	ATOM	3336	OD2	ASP	A	132	-2.735	68.323	-10.142	1.00	26.22	T3
55	ATOM	3337	C	ASP	A	132	-4.248	70.523	-13.290	1.00	41.06	T3
	ATOM	3338	O	ASP	A	132	-5.020	70.359	-14.237	1.00	34.41	T3
	ATOM	3339	N	GLY	A	133	-3.420	71.558	-13.195	1.00	29.66	T3
	ATOM	3340	CA	GLY	A	133	-3.370	72.579	-14.222	1.00	36.13	T3
	ATOM	3341	C	GLY	A	133	-3.150	72.076	-15.636	1.00	30.23	T3
60	ATOM	3342	O	GLY	A	133	-3.650	72.697	-16.569	1.00	27.07	T3
	ATOM	3343	N	ASP	A	134	-2.417	70.974	-15.807	1.00	23.74	T3
	ATOM	3344	CA	ASP	A	134	-2.163	70.443	-17.143	1.00	29.59	T3
	ATOM	3345	CB	ASP	A	134	-1.152	69.305	-17.153	1.00	26.49	T3
	ATOM	3346	CG	ASP	A	134	-0.230	69.339	-16.011	1.00	30.71	T3
65	ATOM	3347	OD1	ASP	A	134	0.526	70.318	-15.931	1.00	36.31	T3
	ATOM	3348	OD2	ASP	A	134	-0.257	68.387	-15.204	1.00	30.68	T3

	ATOM	3349	C	ASP	A	134	-3.398	69.828	-17.733	1.00	27.23	T3
	ATOM	3350	O	ASP	A	134	-3.856	70.199	-18.803	1.00	28.97	T3
	ATOM	3351	N	VAL	A	135	-3.910	68.841	-17.025	1.00	36.76	T3
	ATOM	3352	CA	VAL	A	135	-5.045	68.081	-17.488	1.00	25.87	T3
5	ATOM	3353	CB	VAL	A	135	-5.155	66.818	-16.650	1.00	24.83	T3
	ATOM	3354	CG1	VAL	A	135	-3.873	66.011	-16.817	1.00	23.33	T3
	ATOM	3355	CG2	VAL	A	135	-5.361	67.181	-15.184	1.00	36.44	T3
	ATOM	3356	C	VAL	A	135	-6.420	68.723	-17.630	1.00	30.27	T3
	ATOM	3357	O	VAL	A	135	-7.138	68.383	-18.563	1.00	32.37	T3
10	ATOM	3358	N	THR	A	136	-6.817	69.623	-16.738	1.00	33.74	T3
	ATOM	3359	CA	THR	A	136	-8.140	70.217	-16.904	1.00	32.60	T3
	ATOM	3360	CB	THR	A	136	-9.145	69.684	-15.832	1.00	29.99	T3
	ATOM	3361	OG1	THR	A	136	-8.917	70.327	-14.579	1.00	21.06	T3
	ATOM	3362	CG2	THR	A	136	-8.966	68.193	-15.630	1.00	34.08	T3
15	ATOM	3363	C	THR	A	136	-8.125	71.751	-16.919	1.00	37.22	T3
	ATOM	3364	O	THR	A	136	-7.690	72.405	-15.967	1.00	31.24	T3
	ATOM	3365	N	PHE	A	137	-8.605	72.313	-18.025	1.00	32.07	T3
	ATOM	3366	CA	PHE	A	137	-8.638	73.756	-18.214	1.00	33.69	T3
	ATOM	3367	CB	PHE	A	137	-7.336	74.195	-18.877	1.00	27.08	T3
20	ATOM	3368	CG	PHE	A	137	-6.905	73.312	-20.019	1.00	26.29	T3
	ATOM	3369	CD1	PHE	A	137	-7.458	73.455	-21.280	1.00	26.82	T3
	ATOM	3370	CD2	PHE	A	137	-5.943	72.339	-19.832	1.00	27.61	T3
	ATOM	3371	CE1	PHE	A	137	-7.053	72.643	-22.331	1.00	31.52	T3
	ATOM	3372	CE2	PHE	A	137	-5.538	71.527	-20.878	1.00	31.62	T3
25	ATOM	3373	CZ	PHE	A	137	-6.094	71.681	-22.124	1.00	27.89	T3
	ATOM	3374	C	PHE	A	137	-9.855	74.221	-19.030	1.00	30.06	T3
	ATOM	3375	O	PHE	A	137	-10.513	73.419	-19.690	1.00	32.70	T3
	ATOM	3376	N	PHE	A	138	-10.140	75.520	-18.992	1.00	30.37	T3
	ATOM	3377	CA	PHE	A	138	-11.296	76.064	-19.688	1.00	26.97	T3
30	ATOM	3378	CB	PHE	A	138	-12.334	76.449	-18.629	1.00	27.61	T3
	ATOM	3379	CG	PHE	A	138	-13.706	76.722	-19.174	1.00	25.55	T3
	ATOM	3380	CD1	PHE	A	138	-14.108	76.200	-20.396	1.00	24.64	T3
	ATOM	3381	CD2	PHE	A	138	-14.594	77.520	-18.462	1.00	26.87	T3
	ATOM	3382	CE1	PHE	A	138	-15.370	76.471	-20.901	1.00	31.13	T3
35	ATOM	3383	CE2	PHE	A	138	-15.856	77.798	-18.957	1.00	30.96	T3
	ATOM	3384	CZ	PHE	A	138	-16.245	77.274	-20.180	1.00	24.90	T3
	ATOM	3385	C	PHE	A	138	-10.931	77.238	-20.614	1.00	31.19	T3
	ATOM	3386	O	PHE	A	138	-10.253	78.189	-20.211	1.00	34.09	T3
	ATOM	3387	N	GLY	A	139	-11.426	77.149	-21.851	1.00	32.89	T3
40	ATOM	3388	CA	GLY	A	139	-11.134	78.096	-22.926	1.00	31.21	T3
	ATOM	3389	C	GLY	A	139	-11.618	79.521	-23.069	1.00	22.01	T3
	ATOM	3390	O	GLY	A	139	-11.543	80.287	-22.123	1.00	28.52	T3
	ATOM	3391	N	ALA	A	140	-12.052	79.870	-24.283	1.00	31.93	T3
	ATOM	3392	CA	ALA	A	140	-12.566	81.211	-24.658	1.00	26.19	T3
45	ATOM	3393	CB	ALA	A	140	-13.341	81.831	-23.507	1.00	30.85	T3
	ATOM	3394	C	ALA	A	140	-11.556	82.244	-25.198	1.00	31.93	T3
	ATOM	3395	O	ALA	A	140	-10.785	82.839	-24.447	1.00	32.27	T3
	ATOM	3396	N	LEU	A	141	-11.598	82.462	-26.511	1.00	31.29	T3
	ATOM	3397	CA	LEU	A	141	-10.723	83.411	-27.199	1.00	34.20	T3
50	ATOM	3398	CB	LEU	A	141	-9.561	82.665	-27.871	1.00	29.60	T3
	ATOM	3399	CG	LEU	A	141	-8.620	83.427	-28.811	1.00	33.23	T3
	ATOM	3400	CD1	LEU	A	141	-7.352	82.649	-29.018	1.00	31.81	T3
	ATOM	3401	CD2	LEU	A	141	-9.290	83.656	-30.140	1.00	29.96	T3
	ATOM	3402	C	LEU	A	141	-11.559	84.127	-28.256	1.00	27.04	T3
55	ATOM	3403	O	LEU	A	141	-12.306	83.489	-28.991	1.00	32.61	T3
	ATOM	3404	N	LYS	A	142	-11.429	85.445	-28.350	1.00	29.72	T3
	ATOM	3405	CA	LYS	A	142	-12.212	86.199	-29.322	1.00	32.79	T3
	ATOM	3406	CB	LYS	A	142	-12.501	87.598	-28.787	1.00	24.88	T3
	ATOM	3407	CG	LYS	A	142	-13.334	88.419	-29.744	1.00	28.06	T3
60	ATOM	3408	CD	LYS	A	142	-13.874	89.680	-29.110	1.00	27.47	T3
	ATOM	3409	CE	LYS	A	142	-14.694	90.451	-30.130	1.00	33.18	T3
	ATOM	3410	NZ	LYS	A	142	-15.214	91.723	-29.566	1.00	32.73	T3
	ATOM	3411	C	LYS	A	142	-11.615	86.316	-30.727	1.00	28.68	T3
	ATOM	3412	O	LYS	A	142	-10.471	86.714	-30.892	1.00	35.94	T3
65	ATOM	3413	N	LEU	A	143	-12.409	85.978	-31.738	1.00	30.98	T3
	ATOM	3414	CA	LEU	A	143	-11.968	86.053	-33.127	1.00	32.94	T3

	ATOM	3415	CB	LEU	A	143	-12.818	85.135	-34.001	1.00	38.43	T3
	ATOM	3416	CG	LEU	A	143	-12.880	83.645	-33.654	1.00	31.92	T3
	ATOM	3417	CD1	LEU	A	143	-13.856	82.953	-34.582	1.00	34.25	T3
	ATOM	3418	CD2	LEU	A	143	-11.501	83.022	-33.784	1.00	22.49	T3
5	ATOM	3419	C	LEU	A	143	-12.098	87.474	-33.645	1.00	31.13	T3
	ATOM	3420	O	LEU	A	143	-12.897	88.253	-33.138	1.00	29.98	T3
	ATOM	3421	N	LEU	A	144	-11.313	87.813	-34.659	1.00	26.91	T3
	ATOM	3422	CA	LEU	A	144	-11.369	89.153	-35.235	1.00	32.73	T3
	ATOM	3423	CB	LEU	A	144	-10.016	89.551	-35.835	1.00	27.45	T3
10	ATOM	3424	CG	LEU	A	144	-8.870	89.717	-34.835	1.00	24.32	T3
	ATOM	3425	CD1	LEU	A	144	-7.577	89.912	-35.595	1.00	29.70	T3
	ATOM	3426	CD2	LEU	A	144	-9.132	90.900	-33.904	1.00	26.52	T3
	ATOM	3427	C	LEU	A	144	-12.424	89.206	-36.322	1.00	33.58	T3
	ATOM	3428	O	LEU	A	144	-12.837	88.126	-36.795	1.00	32.13	T3
15	ATOM	3429	OXT	LEU	A	144	-12.811	90.333	-36.695	1.00	33.19	T3
	ATOM	3430	CB	VAL	A	1	14.763	104.864	-24.415	1.00	31.43	T4
	ATOM	3431	CG1	VAL	A	1	13.260	104.617	-24.185	1.00	30.05	T4
	ATOM	3432	CG2	VAL	A	1	14.964	105.866	-25.552	1.00	32.15	T4
	ATOM	3433	C	VAL	A	1	15.323	104.322	-22.006	1.00	26.99	T4
20	ATOM	3434	O	VAL	A	1	14.695	104.561	-20.973	1.00	37.84	T4
	ATOM	3435	N	VAL	A	1	16.864	105.791	-23.335	1.00	30.11	T4
	ATOM	3436	CA	VAL	A	1	15.439	105.398	-23.097	1.00	27.60	T4
	ATOM	3437	N	THR	A	2	15.925	103.148	-22.226	1.00	28.38	T4
	ATOM	3438	CA	THR	A	2	15.844	102.066	-21.238	1.00	33.70	T4
25	ATOM	3439	CB	THR	A	2	15.305	100.766	-21.857	1.00	24.35	T4
	ATOM	3440	OG1	THR	A	2	16.296	100.214	-22.725	1.00	29.12	T4
	ATOM	3441	CG2	THR	A	2	14.029	101.032	-22.648	1.00	25.39	T4
	ATOM	3442	C	THR	A	2	17.187	101.742	-20.601	1.00	29.91	T4
	ATOM	3443	O	THR	A	2	18.223	102.242	-21.032	1.00	31.79	T4
30	ATOM	3444	N	GLN	A	3	17.160	100.887	-19.579	1.00	31.87	T4
	ATOM	3445	CA	GLN	A	3	18.373	100.489	-18.866	1.00	34.19	T4
	ATOM	3446	CB	GLN	A	3	18.206	100.722	-17.363	1.00	25.52	T4
	ATOM	3447	CG	GLN	A	3	17.630	102.073	-16.998	1.00	30.70	T4
	ATOM	3448	CD	GLN	A	3	17.629	102.305	-15.510	1.00	26.99	T4
35	ATOM	3449	OE1	GLN	A	3	18.684	102.325	-14.890	1.00	28.60	T4
	ATOM	3450	NE2	GLN	A	3	16.448	102.482	-14.926	1.00	26.24	T4
	ATOM	3451	C	GLN	A	3	18.710	99.022	-19.092	1.00	28.89	T4
	ATOM	3452	O	GLN	A	3	18.078	98.138	-18.510	1.00	38.16	T4
	ATOM	3453	N	ASP	A	4	19.708	98.754	-19.923	1.00	28.02	T4
40	ATOM	3454	CA	ASP	A	4	20.082	97.376	-20.167	1.00	24.95	T4
	ATOM	3455	CB	ASP	A	4	21.203	97.305	-21.197	1.00	30.48	T4
	ATOM	3456	CG	ASP	A	4	20.780	97.817	-22.557	1.00	32.14	T4
	ATOM	3457	OD1	ASP	A	4	19.557	97.927	-22.788	1.00	25.41	T4
	ATOM	3458	OD2	ASP	A	4	21.663	98.095	-23.404	1.00	33.80	T4
45	ATOM	3459	C	ASP	A	4	20.539	96.716	-18.874	1.00	26.62	T4
	ATOM	3460	O	ASP	A	4	21.097	97.372	-18.001	1.00	27.03	T4
	ATOM	3461	N	CYS	A	5	20.289	95.419	-18.750	1.00	26.47	T4
	ATOM	3462	CA	CYS	A	5	20.690	94.663	-17.567	1.00	37.79	T4
	ATOM	3463	CB	CYS	A	5	19.694	94.865	-16.405	1.00	27.72	T4
50	ATOM	3464	SG	CYS	A	5	17.934	95.177	-16.841	1.00	29.37	T4
	ATOM	3465	C	CYS	A	5	20.795	93.189	-17.914	1.00	27.27	T4
	ATOM	3466	O	CYS	A	5	20.068	92.700	-18.767	1.00	26.31	T4
	ATOM	3467	N	LEU	A	6	21.726	92.490	-17.276	1.00	32.15	T4
	ATOM	3468	CA	LEU	A	6	21.912	91.063	-17.511	1.00	36.12	T4
55	ATOM	3469	CB	LEU	A	6	23.105	90.821	-18.433	1.00	36.37	T4
	ATOM	3470	CG	LEU	A	6	23.455	89.362	-18.728	1.00	32.31	T4
	ATOM	3471	CD1	LEU	A	6	24.091	89.247	-20.095	1.00	30.83	T4
	ATOM	3472	CD2	LEU	A	6	24.388	88.833	-17.661	1.00	32.80	T4
	ATOM	3473	C	LEU	A	6	22.143	90.415	-16.160	1.00	37.97	T4
60	ATOM	3474	O	LEU	A	6	22.914	90.923	-15.364	1.00	35.00	T4
	ATOM	3475	N	GLN	A	7	21.468	89.302	-15.895	1.00	26.19	T4
	ATOM	3476	CA	GLN	A	7	21.605	88.629	-14.617	1.00	27.69	T4
	ATOM	3477	CB	GLN	A	7	20.338	88.817	-13.805	1.00	31.14	T4
	ATOM	3478	CG	GLN	A	7	20.456	88.394	-12.365	1.00	31.64	T4
65	ATOM	3479	CD	GLN	A	7	19.256	88.824	-11.554	1.00	29.44	T4
	ATOM	3480	OE1	GLN	A	7	18.161	88.307	-11.731	1.00	38.75	T4

	ATOM	3481	NE2	GLN	A	7	19.453	89.789	-10.672	1.00	35.81	T4
	ATOM	3482	C	GLN	A	7	21.885	87.153	-14.775	1.00	34.75	T4
	ATOM	3483	O	GLN	A	7	21.320	86.494	-15.642	1.00	31.48	T4
	ATOM	3484	N	LEU	A	8	22.758	86.636	-13.920	1.00	32.76	T4
5	ATOM	3485	CA	LEU	A	8	23.127	85.230	-13.956	1.00	30.73	T4
	ATOM	3486	CB	LEU	A	8	24.637	85.091	-14.153	1.00	24.31	T4
	ATOM	3487	CG	LEU	A	8	25.256	85.103	-15.558	1.00	31.72	T4
	ATOM	3488	CD1	LEU	A	8	24.275	85.588	-16.584	1.00	28.07	T4
	ATOM	3489	CD2	LEU	A	8	26.500	85.971	-15.537	1.00	26.28	T4
10	ATOM	3490	C	LEU	A	8	22.708	84.505	-12.686	1.00	26.94	T4
	ATOM	3491	O	LEU	A	8	22.590	85.108	-11.621	1.00	28.89	T4
	ATOM	3492	N	ILE	A	9	22.498	83.201	-12.816	1.00	31.23	T4
	ATOM	3493	CA	ILE	A	9	22.081	82.342	-11.711	1.00	28.02	T4
	ATOM	3494	CB	ILE	A	9	20.658	81.824	-11.948	1.00	33.28	T4
15	ATOM	3495	CG2	ILE	A	9	20.386	80.619	-11.098	1.00	29.62	T4
	ATOM	3496	CG1	ILE	A	9	19.640	82.893	-11.619	1.00	22.17	T4
	ATOM	3497	CD1	ILE	A	9	18.243	82.410	-11.899	1.00	35.16	T4
	ATOM	3498	C	ILE	A	9	22.996	81.123	-11.632	1.00	31.84	T4
	ATOM	3499	O	ILE	A	9	23.422	80.602	-12.658	1.00	31.44	T4
20	ATOM	3500	N	ALA	A	10	23.282	80.653	-10.426	1.00	33.43	T4
	ATOM	3501	CA	ALA	A	10	24.131	79.481	-10.281	1.00	34.48	T4
	ATOM	3502	CB	ALA	A	10	24.356	79.177	-8.830	1.00	23.96	T4
	ATOM	3503	C	ALA	A	10	23.507	78.272	-10.955	1.00	30.70	T4
	ATOM	3504	O	ALA	A	10	22.327	77.969	-10.747	1.00	35.58	T4
25	ATOM	3505	N	ASP	A	11	24.305	77.577	-11.762	1.00	28.90	T4
	ATOM	3506	CA	ASP	A	11	23.827	76.381	-12.458	1.00	31.28	T4
	ATOM	3507	CB	ASP	A	11	24.494	76.238	-13.825	1.00	26.25	T4
	ATOM	3508	CG	ASP	A	11	24.132	74.947	-14.499	1.00	28.61	T4
	ATOM	3509	OD1	ASP	A	11	22.964	74.536	-14.370	1.00	28.68	T4
30	ATOM	3510	OD2	ASP	A	11	25.005	74.346	-15.156	1.00	30.68	T4
	ATOM	3511	C	ASP	A	11	24.107	75.150	-11.617	1.00	29.60	T4
	ATOM	3512	O	ASP	A	11	25.168	74.543	-11.719	1.00	25.50	T4
	ATOM	3513	N	SER	A	12	23.132	74.806	-10.783	1.00	37.37	T4
	ATOM	3514	CA	SER	A	12	23.214	73.673	-9.871	1.00	31.66	T4
35	ATOM	3515	CB	SER	A	12	21.983	73.646	-8.960	1.00	30.01	T4
	ATOM	3516	OG	SER	A	12	20.775	73.553	-9.714	1.00	25.37	T4
	ATOM	3517	C	SER	A	12	23.331	72.333	-10.569	1.00	22.94	T4
	ATOM	3518	O	SER	A	12	23.001	71.298	-9.991	1.00	25.03	T4
	ATOM	3519	N	GLU	A	13	23.793	72.337	-11.814	1.00	31.02	T4
40	ATOM	3520	CA	GLU	A	13	23.931	71.081	-12.530	1.00	26.44	T4
	ATOM	3521	CB	GLU	A	13	22.741	70.869	-13.439	1.00	27.60	T4
	ATOM	3522	CG	GLU	A	13	21.633	70.140	-12.722	1.00	31.50	T4
	ATOM	3523	CD	GLU	A	13	20.481	69.848	-13.648	1.00	34.15	T4
	ATOM	3524	OE1	GLU	A	13	20.766	69.454	-14.812	1.00	24.85	T4
45	ATOM	3525	OE2	GLU	A	13	19.299	70.003	-13.224	1.00	25.04	T4
	ATOM	3526	C	GLU	A	13	25.213	70.872	-13.303	1.00	30.12	T4
	ATOM	3527	O	GLU	A	13	25.250	70.121	-14.274	1.00	35.18	T4
	ATOM	3528	N	THR	A	14	26.262	71.551	-12.866	1.00	34.54	T4
	ATOM	3529	CA	THR	A	14	27.575	71.402	-13.464	1.00	30.83	T4
50	ATOM	3530	CB	THR	A	14	27.837	72.390	-14.615	1.00	29.18	T4
	ATOM	3531	OG1	THR	A	14	27.675	73.725	-14.139	1.00	40.81	T4
	ATOM	3532	CG2	THR	A	14	26.883	72.137	-15.769	1.00	30.78	T4
	ATOM	3533	C	THR	A	14	28.545	71.663	-12.327	1.00	31.55	T4
	ATOM	3534	O	THR	A	14	28.242	72.399	-11.387	1.00	26.08	T4
55	ATOM	3535	N	PRO	A	15	29.722	71.036	-12.389	1.00	32.64	T4
	ATOM	3536	CD	PRO	A	15	30.180	70.157	-13.479	1.00	30.91	T4
	ATOM	3537	CA	PRO	A	15	30.755	71.181	-11.363	1.00	27.66	T4
	ATOM	3538	CB	PRO	A	15	31.866	70.255	-11.859	1.00	30.35	T4
	ATOM	3539	CG	PRO	A	15	31.142	69.268	-12.760	1.00	25.09	T4
60	ATOM	3540	C	PRO	A	15	31.233	72.615	-11.268	1.00	26.98	T4
	ATOM	3541	O	PRO	A	15	31.286	73.315	-12.275	1.00	31.54	T4
	ATOM	3542	N	THR	A	16	31.585	73.049	-10.065	1.00	31.08	T4
	ATOM	3543	CA	THR	A	16	32.084	74.403	-9.883	1.00	23.49	T4
	ATOM	3544	CB	THR	A	16	32.151	74.750	-8.410	1.00	34.10	T4
65	ATOM	3545	OG1	THR	A	16	33.194	73.992	-7.788	1.00	27.76	T4
	ATOM	3546	CG2	THR	A	16	30.836	74.399	-7.744	1.00	29.70	T4

	ATOM	3547	C	THR	A	16	33.484	74.472	-10.473	1.00	40.93	T4
	ATOM	3548	O	THR	A	16	34.388	73.804	-9.995	1.00	32.84	T4
	ATOM	3549	N	ILE	A	17	33.659	75.276	-11.515	1.00	33.34	T4
	ATOM	3550	CA	ILE	A	17	34.952	75.418	-12.186	1.00	33.81	T4
5	ATOM	3551	CB	ILE	A	17	34.892	76.524	-13.241	1.00	23.23	T4
	ATOM	3552	CG2	ILE	A	17	36.243	76.688	-13.902	1.00	23.89	T4
	ATOM	3553	CG1	ILE	A	17	33.815	76.189	-14.270	1.00	37.92	T4
	ATOM	3554	CD1	ILE	A	17	33.626	77.247	-15.325	1.00	32.96	T4
	ATOM	3555	C	ILE	A	17	36.141	75.713	-11.257	1.00	29.34	T4
10	ATOM	3556	O	ILE	A	17	36.086	76.608	-10.400	1.00	28.62	T4
	ATOM	3557	N	GLN	A	18	37.217	74.946	-11.439	1.00	36.59	T4
	ATOM	3558	CA	GLN	A	18	38.430	75.107	-10.643	1.00	31.17	T4
	ATOM	3559	CB	GLN	A	18	38.819	73.790	-9.999	1.00	29.55	T4
	ATOM	3560	CG	GLN	A	18	38.953	73.915	-8.517	1.00	25.02	T4
15	ATOM	3561	CD	GLN	A	18	37.603	74.042	-7.842	1.00	30.78	T4
	ATOM	3562	OE1	GLN	A	18	36.881	73.053	-7.690	1.00	26.86	T4
	ATOM	3563	NE2	GLN	A	18	37.244	75.262	-7.445	1.00	30.49	T4
	ATOM	3564	C	GLN	A	18	39.581	75.598	-11.510	1.00	30.78	T4
	ATOM	3565	O	GLN	A	18	39.805	75.080	-12.608	1.00	33.98	T4
20	ATOM	3566	N	LYS	A	19	40.320	76.589	-11.021	1.00	30.08	T4
	ATOM	3567	CA	LYS	A	19	41.434	77.129	-11.792	1.00	25.59	T4
	ATOM	3568	CB	LYS	A	19	40.902	77.797	-13.058	1.00	23.61	T4
	ATOM	3569	CG	LYS	A	19	41.935	78.610	-13.802	1.00	28.30	T4
	ATOM	3570	CD	LYS	A	19	41.396	79.070	-15.158	1.00	25.87	T4
25	ATOM	3571	CE	LYS	A	19	42.433	79.922	-15.909	1.00	33.21	T4
	ATOM	3572	NZ	LYS	A	19	41.965	80.343	-17.272	1.00	32.10	T4
	ATOM	3573	C	LYS	A	19	42.276	78.120	-11.000	1.00	35.53	T4
	ATOM	3574	O	LYS	A	19	41.745	79.049	-10.389	1.00	40.45	T4
	ATOM	3575	N	GLY	A	20	43.594	77.922	-11.032	1.00	29.07	T4
30	ATOM	3576	CA	GLY	A	20	44.500	78.796	-10.307	1.00	36.37	T4
	ATOM	3577	C	GLY	A	20	44.107	78.868	-8.849	1.00	27.77	T4
	ATOM	3578	O	GLY	A	20	44.140	79.948	-8.258	1.00	29.16	T4
	ATOM	3579	N	SER	A	21	43.729	77.717	-8.284	1.00	31.67	T4
	ATOM	3580	CA	SER	A	21	43.301	77.610	-6.877	1.00	27.89	T4
35	ATOM	3581	CB	SER	A	21	44.505	77.751	-5.922	1.00	26.25	T4
	ATOM	3582	OG	SER	A	21	45.138	79.018	-6.038	1.00	31.72	T4
	ATOM	3583	C	SER	A	21	42.205	78.631	-6.507	1.00	31.13	T4
	ATOM	3584	O	SER	A	21	42.196	79.207	-5.410	1.00	27.50	T4
	ATOM	3585	N	TYR	A	22	41.292	78.835	-7.456	1.00	35.46	T4
40	ATOM	3586	CA	TYR	A	22	40.154	79.738	-7.320	1.00	33.43	T4
	ATOM	3587	CB	TYR	A	22	40.288	80.916	-8.279	1.00	33.42	T4
	ATOM	3588	CG	TYR	A	22	40.903	82.143	-7.672	1.00	28.57	T4
	ATOM	3589	CD1	TYR	A	22	41.744	82.052	-6.563	1.00	25.29	T4
	ATOM	3590	CE1	TYR	A	22	42.346	83.197	-6.013	1.00	27.48	T4
45	ATOM	3591	CD2	TYR	A	22	40.673	83.401	-8.220	1.00	31.77	T4
	ATOM	3592	CE2	TYR	A	22	41.271	84.550	-7.680	1.00	35.51	T4
	ATOM	3593	CZ	TYR	A	22	42.107	84.436	-6.575	1.00	31.65	T4
	ATOM	3594	OH	TYR	A	22	42.709	85.546	-6.031	1.00	36.85	T4
	ATOM	3595	C	TYR	A	22	38.931	78.936	-7.711	1.00	36.79	T4
50	ATOM	3596	O	TYR	A	22	39.009	78.063	-8.581	1.00	29.58	T4
	ATOM	3597	N	THR	A	23	37.804	79.212	-7.067	1.00	33.40	T4
	ATOM	3598	CA	THR	A	23	36.581	78.504	-7.410	1.00	31.20	T4
	ATOM	3599	CB	THR	A	23	35.844	77.984	-6.170	1.00	36.55	T4
	ATOM	3600	OG1	THR	A	23	36.784	77.460	-5.227	1.00	33.78	T4
55	ATOM	3601	CG2	THR	A	23	34.907	76.870	-6.570	1.00	31.16	T4
	ATOM	3602	C	THR	A	23	35.666	79.461	-8.169	1.00	29.39	T4
	ATOM	3603	O	THR	A	23	35.434	80.592	-7.733	1.00	33.08	T4
	ATOM	3604	N	PHE	A	24	35.161	79.010	-9.313	1.00	30.93	T4
	ATOM	3605	CA	PHE	A	24	34.273	79.829	-10.124	1.00	31.06	T4
60	ATOM	3606	CB	PHE	A	24	34.876	80.056	-11.511	1.00	32.95	T4
	ATOM	3607	CG	PHE	A	24	36.133	80.869	-11.496	1.00	35.43	T4
	ATOM	3608	CD1	PHE	A	24	37.348	80.292	-11.171	1.00	32.98	T4
	ATOM	3609	CD2	PHE	A	24	36.095	82.226	-11.781	1.00	29.74	T4
	ATOM	3610	CE1	PHE	A	24	38.510	81.059	-11.130	1.00	25.37	T4
65	ATOM	3611	CE2	PHE	A	24	37.246	82.998	-11.742	1.00	32.56	T4
	ATOM	3612	CZ	PHE	A	24	38.455	82.412	-11.415	1.00	34.93	T4

	ATOM	3613	C	PHE	A	24	32.893	79.202	-10.258	1.00	26.09	T4
	ATOM	3614	O	PHE	A	24	32.754	78.061	-10.683	1.00	35.80	T4
	ATOM	3615	N	VAL	A	25	31.873	79.964	-9.885	1.00	30.91	T4
	ATOM	3616	CA	VAL	A	25	30.496	79.503	-9.961	1.00	33.17	T4
5	ATOM	3617	CB	VAL	A	25	29.538	80.530	-9.334	1.00	29.66	T4
	ATOM	3618	CG1	VAL	A	25	28.103	80.070	-9.503	1.00	23.79	T4
	ATOM	3619	CG2	VAL	A	25	29.876	80.727	-7.868	1.00	32.46	T4
	ATOM	3620	C	VAL	A	25	30.071	79.283	-11.404	1.00	36.04	T4
	ATOM	3621	O	VAL	A	25	30.335	80.113	-12.272	1.00	28.84	T4
10	ATOM	3622	N	PRO	A	26	29.422	78.148	-11.682	1.00	34.82	T4
	ATOM	3623	CD	PRO	A	26	29.253	76.998	-10.784	1.00	28.25	T4
	ATOM	3624	CA	PRO	A	26	28.953	77.822	-13.031	1.00	28.09	T4
	ATOM	3625	CB	PRO	A	26	28.630	76.332	-12.935	1.00	29.15	T4
	ATOM	3626	CG	PRO	A	26	29.387	75.859	-11.732	1.00	32.22	T4
15	ATOM	3627	C	PRO	A	26	27.689	78.642	-13.264	1.00	33.38	T4
	ATOM	3628	O	PRO	A	26	26.664	78.386	-12.629	1.00	25.35	T4
	ATOM	3629	N	TRP	A	27	27.739	79.619	-14.162	1.00	30.28	T4
	ATOM	3630	CA	TRP	A	27	26.561	80.449	-14.390	1.00	33.23	T4
	ATOM	3631	CB	TRP	A	27	26.972	81.867	-14.796	1.00	30.50	T4
20	ATOM	3632	CG	TRP	A	27	27.730	82.578	-13.744	1.00	32.01	T4
	ATOM	3633	CD2	TRP	A	27	27.327	82.781	-12.389	1.00	34.02	T4
	ATOM	3634	CE2	TRP	A	27	28.382	83.449	-11.732	1.00	29.41	T4
	ATOM	3635	CE3	TRP	A	27	26.179	82.459	-11.663	1.00	26.72	T4
	ATOM	3636	CD1	TRP	A	27	28.976	83.123	-13.858	1.00	35.03	T4
25	ATOM	3637	NE1	TRP	A	27	29.377	83.648	-12.653	1.00	27.93	T4
	ATOM	3638	CZ2	TRP	A	27	28.322	83.801	-10.383	1.00	29.26	T4
	ATOM	3639	CZ3	TRP	A	27	26.118	82.811	-10.315	1.00	28.59	T4
	ATOM	3640	CH2	TRP	A	27	27.186	83.474	-9.694	1.00	34.08	T4
	ATOM	3641	C	TRP	A	27	25.550	79.916	-15.392	1.00	31.53	T4
30	ATOM	3642	O	TRP	A	27	25.825	79.008	-16.178	1.00	29.39	T4
	ATOM	3643	N	LEU	A	28	24.365	80.508	-15.337	1.00	31.62	T4
	ATOM	3644	CA	LEU	A	28	23.260	80.159	-16.216	1.00	33.47	T4
	ATOM	3645	CB	LEU	A	28	22.417	79.069	-15.573	1.00	26.67	T4
	ATOM	3646	CG	LEU	A	28	21.529	78.298	-16.539	1.00	28.89	T4
35	ATOM	3647	CD1	LEU	A	28	22.415	77.568	-17.554	1.00	32.29	T4
	ATOM	3648	CD2	LEU	A	28	20.659	77.319	-15.743	1.00	29.14	T4
	ATOM	3649	C	LEU	A	28	22.441	81.437	-16.368	1.00	27.09	T4
	ATOM	3650	O	LEU	A	28	22.089	82.079	-15.372	1.00	29.16	T4
	ATOM	3651	N	LEU	A	29	22.138	81.814	-17.601	1.00	28.97	T4
40	ATOM	3652	CA	LEU	A	29	21.397	83.042	-17.823	1.00	30.44	T4
	ATOM	3653	CB	LEU	A	29	21.108	83.237	-19.306	1.00	28.93	T4
	ATOM	3654	CG	LEU	A	29	20.307	84.504	-19.581	1.00	27.26	T4
	ATOM	3655	CD1	LEU	A	29	21.206	85.713	-19.454	1.00	28.46	T4
	ATOM	3656	CD2	LEU	A	29	19.708	84.435	-20.948	1.00	39.45	T4
45	ATOM	3657	C	LEU	A	29	20.083	83.088	-17.070	1.00	29.96	T4
	ATOM	3658	O	LEU	A	29	19.260	82.173	-17.180	1.00	25.27	T4
	ATOM	3659	N	SER	A	30	19.892	84.147	-16.290	1.00	29.75	T4
	ATOM	3660	CA	SER	A	30	18.641	84.332	-15.571	1.00	25.82	T4
	ATOM	3661	CB	SER	A	30	18.851	85.168	-14.320	1.00	31.11	T4
50	ATOM	3662	OG	SER	A	30	17.609	85.425	-13.706	1.00	31.31	T4
	ATOM	3663	C	SER	A	30	17.760	85.094	-16.550	1.00	30.03	T4
	ATOM	3664	O	SER	A	30	16.657	84.676	-16.867	1.00	32.36	T4
	ATOM	3665	N	PHE	A	31	18.274	86.216	-17.041	1.00	27.70	T4
	ATOM	3666	CA	PHE	A	31	17.562	87.032	-18.007	1.00	37.01	T4
55	ATOM	3667	CB	PHE	A	31	16.370	87.723	-17.345	1.00	28.23	T4
	ATOM	3668	CG	PHE	A	31	16.703	89.033	-16.693	1.00	32.03	T4
	ATOM	3669	CD1	PHE	A	31	16.737	90.206	-17.442	1.00	28.73	T4
	ATOM	3670	CD2	PHE	A	31	16.979	89.097	-15.335	1.00	25.49	T4
	ATOM	3671	CE1	PHE	A	31	17.039	91.423	-16.849	1.00	27.09	T4
60	ATOM	3672	CE2	PHE	A	31	17.282	90.306	-14.731	1.00	24.91	T4
	ATOM	3673	CZ	PHE	A	31	17.312	91.475	-15.491	1.00	27.42	T4
	ATOM	3674	C	PHE	A	31	18.522	88.063	-18.580	1.00	26.23	T4
	ATOM	3675	O	PHE	A	31	19.497	88.439	-17.941	1.00	31.40	T4
	ATOM	3676	N	LYS	A	32	18.249	88.508	-19.794	1.00	29.44	T4
65	ATOM	3677	CA	LYS	A	32	19.084	89.496	-20.446	1.00	31.07	T4
	ATOM	3678	CB	LYS	A	32	19.948	88.836	-21.512	1.00	25.62	T4

	ATOM	3679	CG	LYS	A	32	20.600	89.833	-22.429	1.00	27.40	T4
	ATOM	3680	CD	LYS	A	32	21.276	89.184	-23.601	1.00	30.05	T4
	ATOM	3681	CE	LYS	A	32	21.662	90.255	-24.599	1.00	35.82	T4
	ATOM	3682	NZ	LYS	A	32	22.494	89.697	-25.698	1.00	33.46	T4
5	ATOM	3683	C	LYS	A	32	18.181	90.536	-21.088	1.00	27.81	T4
	ATOM	3684	O	LYS	A	32	17.293	90.201	-21.869	1.00	30.78	T4
	ATOM	3685	N	ARG	A	33	18.412	91.799	-20.767	1.00	34.66	T4
	ATOM	3686	CA	ARG	A	33	17.595	92.878	-21.300	1.00	34.70	T4
	ATOM	3687	CB	ARG	A	33	16.761	93.468	-20.159	1.00	30.51	T4
10	ATOM	3688	CG	ARG	A	33	15.967	94.717	-20.471	1.00	32.58	T4
	ATOM	3689	CD	ARG	A	33	14.859	94.858	-19.445	1.00	38.80	T4
	ATOM	3690	NE	ARG	A	33	14.082	96.079	-19.608	1.00	29.62	T4
	ATOM	3691	CZ	ARG	A	33	14.461	97.261	-19.148	1.00	24.90	T4
	ATOM	3692	NH1	ARG	A	33	15.609	97.368	-18.492	1.00	35.02	T4
15	ATOM	3693	NH2	ARG	A	33	13.700	98.330	-19.346	1.00	32.47	T4
	ATOM	3694	C	ARG	A	33	18.475	93.940	-21.935	1.00	28.37	T4
	ATOM	3695	O	ARG	A	33	19.377	94.482	-21.281	1.00	27.98	T4
	ATOM	3696	N	GLY	A	34	18.226	94.227	-23.210	1.00	23.53	T4
	ATOM	3697	CA	GLY	A	34	19.016	95.236	-23.887	1.00	24.01	T4
20	ATOM	3698	C	GLY	A	34	20.204	94.677	-24.637	1.00	33.75	T4
	ATOM	3699	O	GLY	A	34	20.309	93.472	-24.865	1.00	30.83	T4
	ATOM	3700	N	SER	A	35	21.122	95.560	-24.999	1.00	33.85	T4
	ATOM	3701	CA	SER	A	35	22.293	95.159	-25.764	1.00	32.94	T4
	ATOM	3702	CB	SER	A	35	22.272	95.871	-27.108	1.00	29.29	T4
25	ATOM	3703	OG	SER	A	35	22.197	97.278	-26.907	1.00	28.83	T4
	ATOM	3704	C	SER	A	35	23.648	95.425	-25.108	1.00	28.18	T4
	ATOM	3705	O	SER	A	35	24.644	94.812	-25.489	1.00	27.83	T4
	ATOM	3706	N	ALA	A	36	23.695	96.329	-24.135	1.00	26.67	T4
	ATOM	3707	CA	ALA	A	36	24.956	96.671	-23.498	1.00	24.81	T4
30	ATOM	3708	CB	ALA	A	36	24.746	97.845	-22.565	1.00	24.12	T4
	ATOM	3709	C	ALA	A	36	25.688	95.548	-22.766	1.00	33.10	T4
	ATOM	3710	O	ALA	A	36	26.878	95.683	-22.488	1.00	32.30	T4
	ATOM	3711	N	LEU	A	37	25.004	94.447	-22.458	1.00	28.17	T4
	ATOM	3712	CA	LEU	A	37	25.646	93.338	-21.746	1.00	31.81	T4
35	ATOM	3713	CB	LEU	A	37	25.230	93.368	-20.279	1.00	33.02	T4
	ATOM	3714	CG	LEU	A	37	25.617	94.651	-19.549	1.00	34.44	T4
	ATOM	3715	CD1	LEU	A	37	24.802	94.791	-18.292	1.00	20.26	T4
	ATOM	3716	CD2	LEU	A	37	27.098	94.635	-19.241	1.00	35.30	T4
	ATOM	3717	C	LEU	A	37	25.335	91.969	-22.349	1.00	29.86	T4
40	ATOM	3718	O	LEU	A	37	24.254	91.755	-22.877	1.00	26.59	T4
	ATOM	3719	N	GLU	A	38	26.287	91.045	-22.256	1.00	37.04	T4
	ATOM	3720	CA	GLU	A	38	26.142	89.695	-22.807	1.00	30.89	T4
	ATOM	3721	CB	GLU	A	38	26.737	89.635	-24.211	1.00	23.46	T4
	ATOM	3722	CG	GLU	A	38	25.849	90.132	-25.329	1.00	30.12	T4
45	ATOM	3723	CD	GLU	A	38	26.604	90.276	-26.651	1.00	34.81	T4
	ATOM	3724	OE1	GLU	A	38	27.507	89.453	-26.924	1.00	26.49	T4
	ATOM	3725	OE2	GLU	A	38	26.284	91.210	-27.421	1.00	30.76	T4
	ATOM	3726	C	GLU	A	38	26.861	88.647	-21.974	1.00	27.68	T4
	ATOM	3727	O	GLU	A	38	27.712	88.978	-21.160	1.00	32.46	T4
50	ATOM	3728	N	GLU	A	39	26.527	87.378	-22.187	1.00	30.57	T4
	ATOM	3729	CA	GLU	A	39	27.207	86.299	-21.478	1.00	32.70	T4
	ATOM	3730	CB	GLU	A	39	26.346	85.066	-21.325	1.00	34.57	T4
	ATOM	3731	CG	GLU	A	39	24.940	85.295	-20.935	1.00	29.54	T4
	ATOM	3732	CD	GLU	A	39	24.056	84.195	-21.494	1.00	30.97	T4
55	ATOM	3733	OE1	GLU	A	39	23.457	84.411	-22.584	1.00	29.96	T4
	ATOM	3734	OE2	GLU	A	39	23.983	83.108	-20.863	1.00	30.62	T4
	ATOM	3735	C	GLU	A	39	28.339	85.883	-22.386	1.00	32.74	T4
	ATOM	3736	O	GLU	A	39	28.204	85.915	-23.611	1.00	31.71	T4
	ATOM	3737	N	LYS	A	40	29.452	85.475	-21.798	1.00	25.26	T4
60	ATOM	3738	CA	LYS	A	40	30.565	85.025	-22.604	1.00	28.18	T4
	ATOM	3739	CB	LYS	A	40	31.324	86.205	-23.202	1.00	30.31	T4
	ATOM	3740	CG	LYS	A	40	32.471	85.772	-24.109	1.00	26.31	T4
	ATOM	3741	CD	LYS	A	40	33.433	86.920	-24.374	1.00	32.10	T4
	ATOM	3742	CE	LYS	A	40	34.689	86.436	-25.085	1.00	33.67	T4
65	ATOM	3743	NZ	LYS	A	40	35.696	87.539	-25.200	1.00	36.72	T4
	ATOM	3744	C	LYS	A	40	31.501	84.187	-21.767	1.00	33.60	T4

	ATOM	3745	O	LYS	A	40	32.249	84.702	-20.948	1.00	27.14	T4
	ATOM	3746	N	GLU	A	41	31.431	82.880	-21.970	1.00	27.73	T4
	ATOM	3747	CA	GLU	A	41	32.283	81.950	-21.257	1.00	33.56	T4
	ATOM	3748	CB	GLU	A	41	33.701	82.059	-21.809	1.00	34.68	T4
5	ATOM	3749	CG	GLU	A	41	33.709	81.942	-23.319	1.00	36.05	T4
	ATOM	3750	CD	GLU	A	41	35.003	82.406	-23.953	1.00	30.13	T4
	ATOM	3751	OE1	GLU	A	41	35.455	83.545	-23.654	1.00	28.96	T4
	ATOM	3752	OE2	GLU	A	41	35.561	81.632	-24.768	1.00	33.55	T4
	ATOM	3753	C	GLU	A	41	32.259	82.209	-19.763	1.00	30.42	T4
10	ATOM	3754	O	GLU	A	41	33.282	82.506	-19.157	1.00	28.88	T4
	ATOM	3755	N	ASN	A	42	31.070	82.116	-19.184	1.00	30.79	T4
	ATOM	3756	CA	ASN	A	42	30.880	82.297	-17.753	1.00	33.69	T4
	ATOM	3757	CB	ASN	A	42	31.657	81.235	-16.995	1.00	36.88	T4
	ATOM	3758	CG	ASN	A	42	30.869	80.659	-15.868	1.00	33.27	T4
15	ATOM	3759	OD1	ASN	A	42	31.325	80.636	-14.724	1.00	35.92	T4
	ATOM	3760	ND2	ASN	A	42	29.664	80.182	-16.176	1.00	36.28	T4
	ATOM	3761	C	ASN	A	42	31.247	83.662	-17.201	1.00	28.54	T4
	ATOM	3762	O	ASN	A	42	31.474	83.807	-16.003	1.00	38.14	T4
	ATOM	3763	N	LYS	A	43	31.301	84.658	-18.074	1.00	37.86	T4
20	ATOM	3764	CA	LYS	A	43	31.622	86.015	-17.664	1.00	27.02	T4
	ATOM	3765	CB	LYS	A	43	33.041	86.375	-18.094	1.00	31.32	T4
	ATOM	3766	CG	LYS	A	43	34.121	85.628	-17.347	1.00	32.09	T4
	ATOM	3767	CD	LYS	A	43	35.499	85.986	-17.862	1.00	37.59	T4
	ATOM	3768	CE	LYS	A	43	35.724	85.393	-19.243	1.00	24.67	T4
25	ATOM	3769	NZ	LYS	A	43	37.083	85.693	-19.791	1.00	29.81	T4
	ATOM	3770	C	LYS	A	43	30.643	86.970	-18.320	1.00	31.44	T4
	ATOM	3771	O	LYS	A	43	29.996	86.620	-19.306	1.00	25.65	T4
	ATOM	3772	N	ILE	A	44	30.522	88.172	-17.775	1.00	25.19	T4
	ATOM	3773	CA	ILE	A	44	29.632	89.160	-18.357	1.00	31.59	T4
30	ATOM	3774	CB	ILE	A	44	28.948	89.989	-17.280	1.00	30.83	T4
	ATOM	3775	CG2	ILE	A	44	28.040	91.004	-17.920	1.00	33.13	T4
	ATOM	3776	CG1	ILE	A	44	28.149	89.074	-16.364	1.00	25.67	T4
	ATOM	3777	CD1	ILE	A	44	27.433	89.789	-15.266	1.00	28.06	T4
	ATOM	3778	C	ILE	A	44	30.461	90.070	-19.247	1.00	33.28	T4
35	ATOM	3779	O	ILE	A	44	31.437	90.663	-18.803	1.00	34.01	T4
	ATOM	3780	N	LEU	A	45	30.077	90.170	-20.509	1.00	35.40	T4
	ATOM	3781	CA	LEU	A	45	30.803	90.995	-21.457	1.00	25.09	T4
	ATOM	3782	CB	LEU	A	45	30.898	90.264	-22.790	1.00	42.48	T4
	ATOM	3783	CG	LEU	A	45	31.560	91.081	-23.893	1.00	29.49	T4
40	ATOM	3784	CD1	LEU	A	45	33.048	91.235	-23.596	1.00	28.57	T4
	ATOM	3785	CD2	LEU	A	45	31.331	90.391	-25.225	1.00	27.42	T4
	ATOM	3786	C	LEU	A	45	30.167	92.366	-21.675	1.00	34.35	T4
	ATOM	3787	O	LEU	A	45	28.977	92.472	-21.953	1.00	27.46	T4
	ATOM	3788	N	VAL	A	46	30.969	93.417	-21.561	1.00	28.17	T4
45	ATOM	3789	CA	VAL	A	46	30.471	94.776	-21.751	1.00	31.20	T4
	ATOM	3790	CB	VAL	A	46	31.330	95.781	-20.974	1.00	29.11	T4
	ATOM	3791	CG1	VAL	A	46	30.812	97.185	-21.192	1.00	34.37	T4
	ATOM	3792	CG2	VAL	A	46	31.314	95.438	-19.516	1.00	35.56	T4
	ATOM	3793	C	VAL	A	46	30.476	95.162	-23.233	1.00	30.60	T4
50	ATOM	3794	O	VAL	A	46	31.526	95.173	-23.872	1.00	29.76	T4
	ATOM	3795	N	LYS	A	47	29.306	95.489	-23.775	1.00	23.82	T4
	ATOM	3796	CA	LYS	A	47	29.209	95.853	-25.182	1.00	31.36	T4
	ATOM	3797	CB	LYS	A	47	28.023	95.140	-25.828	1.00	28.72	T4
	ATOM	3798	CG	LYS	A	47	28.253	93.671	-26.121	1.00	30.42	T4
55	ATOM	3799	CD	LYS	A	47	29.505	93.504	-26.953	1.00	29.07	T4
	ATOM	3800	CE	LYS	A	47	29.642	92.105	-27.526	1.00	23.72	T4
	ATOM	3801	NZ	LYS	A	47	28.687	91.864	-28.650	1.00	31.35	T4
	ATOM	3802	C	LYS	A	47	29.105	97.344	-25.453	1.00	32.96	T4
	ATOM	3803	O	LYS	A	47	29.177	97.770	-26.597	1.00	30.66	T4
60	ATOM	3804	N	GLU	A	48	28.928	98.133	-24.403	1.00	32.79	T4
	ATOM	3805	CA	GLU	A	48	28.816	99.583	-24.531	1.00	23.98	T4
	ATOM	3806	CB	GLU	A	48	27.361	100.018	-24.535	1.00	37.78	T4
	ATOM	3807	CG	GLU	A	48	26.581	99.649	-25.759	1.00	25.60	T4
	ATOM	3808	CD	GLU	A	48	25.109	99.998	-25.601	1.00	27.19	T4
65	ATOM	3809	OE1	GLU	A	48	24.808	101.090	-25.036	1.00	30.76	T4
	ATOM	3810	OE2	GLU	A	48	24.258	99.185	-26.046	1.00	32.46	T4

	ATOM	3811	C	GLU	A	48	29.464	100.189	-23.314	1.00	34.60	T4
	ATOM	3812	O	GLU	A	48	29.104	99.848	-22.189	1.00	30.64	T4
	ATOM	3813	N	THR	A	49	30.406	101.096	-23.513	1.00	24.80	T4
	ATOM	3814	CA	THR	A	49	31.050	101.690	-22.358	1.00	31.43	T4
5	ATOM	3815	CB	THR	A	49	32.299	102.484	-22.753	1.00	31.03	T4
	ATOM	3816	OG1	THR	A	49	31.939	103.839	-22.997	1.00	35.59	T4
	ATOM	3817	CG2	THR	A	49	32.920	101.898	-24.009	1.00	33.59	T4
	ATOM	3818	C	THR	A	49	30.058	102.603	-21.642	1.00	34.46	T4
	ATOM	3819	O	THR	A	49	29.159	103.158	-22.260	1.00	30.75	T4
10	ATOM	3820	N	GLY	A	50	30.219	102.726	-20.330	1.00	24.12	T4
	ATOM	3821	CA	GLY	A	50	29.343	103.562	-19.532	1.00	27.79	T4
	ATOM	3822	C	GLY	A	50	29.475	103.219	-18.058	1.00	36.98	T4
	ATOM	3823	O	GLY	A	50	30.427	102.556	-17.655	1.00	29.59	T4
	ATOM	3824	N	TYR	A	51	28.519	103.673	-17.252	1.00	33.78	T4
15	ATOM	3825	CA	TYR	A	51	28.519	103.402	-15.818	1.00	30.57	T4
	ATOM	3826	CB	TYR	A	51	28.151	104.665	-15.042	1.00	28.81	T4
	ATOM	3827	CG	TYR	A	51	29.274	105.672	-15.022	1.00	31.12	T4
	ATOM	3828	CD1	TYR	A	51	29.529	106.491	-16.122	1.00	28.48	T4
	ATOM	3829	CE1	TYR	A	51	30.641	107.329	-16.146	1.00	30.94	T4
20	ATOM	3830	CD2	TYR	A	51	30.154	105.727	-13.943	1.00	29.97	T4
	ATOM	3831	CE2	TYR	A	51	31.262	106.552	-13.955	1.00	33.97	T4
	ATOM	3832	CZ	TYR	A	51	31.506	107.345	-15.055	1.00	30.72	T4
	ATOM	3833	OH	TYR	A	51	32.648	108.115	-15.072	1.00	25.02	T4
	ATOM	3834	C	TYR	A	51	27.553	102.281	-15.480	1.00	32.00	T4
25	ATOM	3835	O	TYR	A	51	26.385	102.316	-15.861	1.00	32.15	T4
	ATOM	3836	N	PHE	A	52	28.048	101.278	-14.765	1.00	28.24	T4
	ATOM	3837	CA	PHE	A	52	27.213	100.146	-14.398	1.00	34.94	T4
	ATOM	3838	CB	PHE	A	52	27.707	98.877	-15.089	1.00	32.03	T4
	ATOM	3839	CG	PHE	A	52	27.749	98.969	-16.583	1.00	19.63	T4
30	ATOM	3840	CD1	PHE	A	52	28.743	99.701	-17.222	1.00	28.72	T4
	ATOM	3841	CD2	PHE	A	52	26.805	98.298	-17.358	1.00	32.20	T4
	ATOM	3842	CE1	PHE	A	52	28.797	99.761	-18.615	1.00	31.74	T4
	ATOM	3843	CE2	PHE	A	52	26.851	98.351	-18.743	1.00	34.55	T4
	ATOM	3844	CZ	PHE	A	52	27.849	99.084	-19.375	1.00	27.74	T4
35	ATOM	3845	C	PHE	A	52	27.144	99.869	-12.902	1.00	33.51	T4
	ATOM	3846	O	PHE	A	52	28.094	100.119	-12.161	1.00	34.39	T4
	ATOM	3847	N	PHE	A	53	25.999	99.353	-12.474	1.00	29.91	T4
	ATOM	3848	CA	PHE	A	53	25.782	98.966	-11.094	1.00	29.62	T4
	ATOM	3849	CB	PHE	A	53	24.346	99.231	-10.678	1.00	34.82	T4
40	ATOM	3850	CG	PHE	A	53	23.992	98.658	-9.341	1.00	27.45	T4
	ATOM	3851	CD1	PHE	A	53	24.550	99.171	-8.180	1.00	34.01	T4
	ATOM	3852	CD2	PHE	A	53	23.098	97.596	-9.241	1.00	30.62	T4
	ATOM	3853	CE1	PHE	A	53	24.222	98.638	-6.938	1.00	27.52	T4
	ATOM	3854	CE2	PHE	A	53	22.763	97.056	-8.007	1.00	30.70	T4
45	ATOM	3855	CZ	PHE	A	53	23.326	97.577	-6.852	1.00	29.12	T4
	ATOM	3856	C	PHE	A	53	26.019	97.473	-11.153	1.00	34.46	T4
	ATOM	3857	O	PHE	A	53	25.376	96.783	-11.929	1.00	32.41	T4
	ATOM	3858	N	ILE	A	54	26.943	96.967	-10.349	1.00	30.55	T4
	ATOM	3859	CA	ILE	A	54	27.254	95.545	-10.372	1.00	24.72	T4
50	ATOM	3860	CB	ILE	A	54	28.701	95.346	-10.791	1.00	28.64	T4
	ATOM	3861	CG2	ILE	A	54	28.980	93.879	-11.028	1.00	33.81	T4
	ATOM	3862	CG1	ILE	A	54	28.958	96.148	-12.064	1.00	39.35	T4
	ATOM	3863	CD1	ILE	A	54	30.403	96.415	-12.342	1.00	23.51	T4
	ATOM	3864	C	ILE	A	54	27.024	94.904	-9.016	1.00	31.35	T4
55	ATOM	3865	O	ILE	A	54	27.487	95.415	-8.003	1.00	30.18	T4
	ATOM	3866	N	TYR	A	55	26.309	93.785	-8.996	1.00	34.54	T4
	ATOM	3867	CA	TYR	A	55	26.012	93.108	-7.741	1.00	34.33	T4
	ATOM	3868	CB	TYR	A	55	24.549	93.345	-7.351	1.00	37.98	T4
	ATOM	3869	CG	TYR	A	55	23.550	92.917	-8.401	1.00	27.19	T4
60	ATOM	3870	CD1	TYR	A	55	22.944	91.665	-8.345	1.00	30.38	T4
	ATOM	3871	CE1	TYR	A	55	22.066	91.251	-9.328	1.00	37.30	T4
	ATOM	3872	CD2	TYR	A	55	23.243	93.747	-9.472	1.00	38.00	T4
	ATOM	3873	CE2	TYR	A	55	22.366	93.341	-10.460	1.00	25.43	T4
	ATOM	3874	CZ	TYR	A	55	21.783	92.092	-10.385	1.00	27.83	T4
65	ATOM	3875	OH	TYR	A	55	20.931	91.678	-11.380	1.00	29.77	T4
	ATOM	3876	C	TYR	A	55	26.293	91.622	-7.811	1.00	29.71	T4

	ATOM	3877	O	TYR	A	55	26.593	91.090	-8.866	1.00	28.93	T4
	ATOM	3878	N	GLY	A	56	26.199	90.955	-6.672	1.00	22.97	T4
	ATOM	3879	CA	GLY	A	56	26.458	89.532	-6.643	1.00	24.34	T4
	ATOM	3880	C	GLY	A	56	26.380	88.972	-5.237	1.00	28.44	T4
5	ATOM	3881	O	GLY	A	56	26.838	89.601	-4.284	1.00	31.54	T4
	ATOM	3882	N	GLN	A	57	25.787	87.789	-5.115	1.00	32.66	T4
	ATOM	3883	CA	GLN	A	57	25.646	87.115	-3.835	1.00	32.00	T4
	ATOM	3884	CB	GLN	A	57	24.208	87.225	-3.319	1.00	39.84	T4
	ATOM	3885	CG	GLN	A	57	23.954	86.401	-2.063	1.00	32.75	T4
10	ATOM	3886	CD	GLN	A	57	22.577	86.595	-1.472	1.00	30.81	T4
	ATOM	3887	OE1	GLN	A	57	22.263	87.650	-0.929	1.00	36.28	T4
	ATOM	3888	NE2	GLN	A	57	21.746	85.569	-1.572	1.00	25.78	T4
	ATOM	3889	C	GLN	A	57	26.016	85.646	-3.975	1.00	29.68	T4
	ATOM	3890	O	GLN	A	57	25.800	85.043	-5.019	1.00	29.11	T4
15	ATOM	3891	N	VAL	A	58	26.576	85.086	-2.908	1.00	25.87	T4
	ATOM	3892	CA	VAL	A	58	26.986	83.686	-2.860	1.00	32.42	T4
	ATOM	3893	CB	VAL	A	58	28.521	83.537	-3.065	1.00	29.50	T4
	ATOM	3894	CG1	VAL	A	58	28.959	82.119	-2.755	1.00	31.61	T4
	ATOM	3895	CG2	VAL	A	58	28.893	83.887	-4.483	1.00	28.99	T4
20	ATOM	3896	C	VAL	A	58	26.628	83.120	-1.487	1.00	32.43	T4
	ATOM	3897	O	VAL	A	58	26.775	83.806	-0.481	1.00	26.06	T4
	ATOM	3898	N	LEU	A	59	26.151	81.879	-1.446	1.00	31.28	T4
	ATOM	3899	CA	LEU	A	59	25.812	81.239	-0.177	1.00	35.61	T4
	ATOM	3900	CB	LEU	A	59	24.516	80.436	-0.301	1.00	32.73	T4
25	ATOM	3901	CG	LEU	A	59	23.733	80.187	0.998	1.00	34.22	T4
	ATOM	3902	CD1	LEU	A	59	22.762	79.057	0.785	1.00	33.82	T4
	ATOM	3903	CD2	LEU	A	59	24.661	79.839	2.132	1.00	25.81	T4
	ATOM	3904	C	LEU	A	59	26.955	80.291	0.197	1.00	28.23	T4
	ATOM	3905	O	LEU	A	59	27.195	79.291	-0.489	1.00	33.41	T4
30	ATOM	3906	N	TYR	A	60	27.655	80.598	1.284	1.00	35.31	T4
	ATOM	3907	CA	TYR	A	60	28.770	79.764	1.705	1.00	30.92	T4
	ATOM	3908	CB	TYR	A	60	29.870	80.633	2.312	1.00	23.98	T4
	ATOM	3909	CG	TYR	A	60	30.397	81.633	1.336	1.00	31.10	T4
	ATOM	3910	CD1	TYR	A	60	30.074	82.974	1.445	1.00	37.43	T4
35	ATOM	3911	CE1	TYR	A	60	30.492	83.885	0.495	1.00	31.38	T4
	ATOM	3912	CD2	TYR	A	60	31.160	81.221	0.251	1.00	25.85	T4
	ATOM	3913	CE2	TYR	A	60	31.582	82.111	-0.704	1.00	38.71	T4
	ATOM	3914	CZ	TYR	A	60	31.243	83.443	-0.580	1.00	26.37	T4
	ATOM	3915	OH	TYR	A	60	31.640	84.332	-1.551	1.00	29.64	T4
40	ATOM	3916	C	TYR	A	60	28.382	78.680	2.691	1.00	33.70	T4
	ATOM	3917	O	TYR	A	60	27.776	78.951	3.726	1.00	20.90	T4
	ATOM	3918	N	THR	A	61	28.738	77.445	2.370	1.00	29.78	T4
	ATOM	3919	CA	THR	A	61	28.437	76.323	3.246	1.00	24.04	T4
	ATOM	3920	CB	THR	A	61	27.523	75.312	2.549	1.00	28.48	T4
45	ATOM	3921	OG1	THR	A	61	28.100	74.925	1.293	1.00	30.12	T4
	ATOM	3922	CG2	THR	A	61	26.164	75.930	2.305	1.00	30.01	T4
	ATOM	3923	C	THR	A	61	29.744	75.659	3.626	1.00	33.62	T4
	ATOM	3924	O	THR	A	61	29.768	74.537	4.113	1.00	30.67	T4
	ATOM	3925	N	ASP	A	62	30.832	76.378	3.385	1.00	27.27	T4
50	ATOM	3926	CA	ASP	A	62	32.176	75.911	3.698	1.00	30.55	T4
	ATOM	3927	CB	ASP	A	62	33.164	76.610	2.764	1.00	29.66	T4
	ATOM	3928	CG	ASP	A	62	34.535	75.987	2.785	1.00	28.02	T4
	ATOM	3929	OD1	ASP	A	62	35.061	75.699	1.679	1.00	31.54	T4
	ATOM	3930	OD2	ASP	A	62	35.081	75.795	3.897	1.00	34.71	T4
55	ATOM	3931	C	ASP	A	62	32.448	76.285	5.159	1.00	33.82	T4
	ATOM	3932	O	ASP	A	62	31.929	77.280	5.644	1.00	27.60	T4
	ATOM	3933	N	LYS	A	63	33.245	75.504	5.870	1.00	39.21	T4
	ATOM	3934	CA	LYS	A	63	33.504	75.841	7.261	1.00	31.85	T4
	ATOM	3935	CB	LYS	A	63	33.245	74.628	8.156	1.00	35.19	T4
60	ATOM	3936	CG	LYS	A	63	34.108	73.417	7.830	1.00	29.48	T4
	ATOM	3937	CD	LYS	A	63	33.821	72.249	8.786	1.00	26.92	T4
	ATOM	3938	CE	LYS	A	63	32.349	71.782	8.706	1.00	32.20	T4
	ATOM	3939	NZ	LYS	A	63	32.029	70.649	9.649	1.00	23.42	T4
	ATOM	3940	C	LYS	A	63	34.907	76.368	7.533	1.00	31.56	T4
65	ATOM	3941	O	LYS	A	63	35.377	76.324	8.673	1.00	31.04	T4
	ATOM	3942	N	THR	A	64	35.575	76.893	6.515	1.00	29.51	T4

	ATOM	3943	CA	THR	A	64	36.929	77.372	6.741	1.00	32.94	T4
	ATOM	3944	CB	THR	A	64	37.806	77.232	5.465	1.00	28.19	T4
	ATOM	3945	OG1	THR	A	64	37.140	77.827	4.345	1.00	26.06	T4
	ATOM	3946	CG2	THR	A	64	38.093	75.756	5.179	1.00	31.61	T4
5	ATOM	3947	C	THR	A	64	37.077	78.784	7.302	1.00	29.52	T4
	ATOM	3948	O	THR	A	64	37.682	79.648	6.677	1.00	33.71	T4
	ATOM	3949	N	TYR	A	65	36.525	79.000	8.493	1.00	32.26	T4
	ATOM	3950	CA	TYR	A	65	36.607	80.275	9.222	1.00	27.92	T4
	ATOM	3951	CB	TYR	A	65	37.965	80.361	9.939	1.00	35.20	T4
10	ATOM	3952	CG	TYR	A	65	38.984	81.253	9.274	1.00	25.73	T4
	ATOM	3953	CD1	TYR	A	65	39.098	82.602	9.623	1.00	19.69	T4
	ATOM	3954	CE1	TYR	A	65	40.020	83.446	8.986	1.00	37.78	T4
	ATOM	3955	CD2	TYR	A	65	39.816	80.759	8.273	1.00	33.58	T4
	ATOM	3956	CE2	TYR	A	65	40.744	81.589	7.622	1.00	25.41	T4
15	ATOM	3957	CZ	TYR	A	65	40.841	82.932	7.982	1.00	33.07	T4
	ATOM	3958	OH	TYR	A	65	41.746	83.745	7.320	1.00	26.35	T4
	ATOM	3959	C	TYR	A	65	36.343	81.594	8.487	1.00	33.43	T4
	ATOM	3960	O	TYR	A	65	35.985	82.589	9.120	1.00	28.88	T4
	ATOM	3961	N	ALA	A	66	36.520	81.627	7.172	1.00	29.09	T4
20	ATOM	3962	CA	ALA	A	66	36.291	82.860	6.433	1.00	35.99	T4
	ATOM	3963	CB	ALA	A	66	37.431	83.817	6.674	1.00	29.81	T4
	ATOM	3964	C	ALA	A	66	36.137	82.610	4.945	1.00	31.46	T4
	ATOM	3965	O	ALA	A	66	37.058	82.121	4.287	1.00	28.50	T4
	ATOM	3966	N	MET	A	67	34.965	82.943	4.417	1.00	31.04	T4
25	ATOM	3967	CA	MET	A	67	34.699	82.763	2.999	1.00	31.21	T4
	ATOM	3968	CB	MET	A	67	33.542	81.783	2.793	1.00	28.57	T4
	ATOM	3969	CG	MET	A	67	33.868	80.354	3.154	1.00	42.76	T4
	ATOM	3970	SD	MET	A	67	35.215	79.758	2.152	1.00	23.31	T4
	ATOM	3971	CE	MET	A	67	34.399	79.316	0.623	1.00	36.99	T4
30	ATOM	3972	C	MET	A	67	34.354	84.112	2.395	1.00	34.35	T4
	ATOM	3973	O	MET	A	67	34.129	85.080	3.123	1.00	33.30	T4
	ATOM	3974	N	GLY	A	68	34.323	84.175	1.067	1.00	28.34	T4
	ATOM	3975	CA	GLY	A	68	34.001	85.419	0.392	1.00	29.53	T4
	ATOM	3976	C	GLY	A	68	34.261	85.355	-1.098	1.00	29.11	T4
35	ATOM	3977	O	GLY	A	68	34.857	84.397	-1.585	1.00	31.43	T4
	ATOM	3978	N	HIS	A	69	33.801	86.362	-1.835	1.00	29.37	T4
	ATOM	3979	CA	HIS	A	69	34.022	86.395	-3.275	1.00	33.73	T4
	ATOM	3980	CB	HIS	A	69	32.774	85.934	-4.047	1.00	30.95	T4
	ATOM	3981	CG	HIS	A	69	31.520	86.671	-3.694	1.00	29.88	T4
40	ATOM	3982	CD2	HIS	A	69	30.837	87.634	-4.357	1.00	26.95	T4
	ATOM	3983	ND1	HIS	A	69	30.803	86.420	-2.543	1.00	38.28	T4
	ATOM	3984	CE1	HIS	A	69	29.734	87.193	-2.514	1.00	35.79	T4
	ATOM	3985	NE2	HIS	A	69	29.730	87.940	-3.603	1.00	27.47	T4
	ATOM	3986	C	HIS	A	69	34.463	87.764	-3.768	1.00	39.65	T4
45	ATOM	3987	O	HIS	A	69	34.398	88.753	-3.040	1.00	27.39	T4
	ATOM	3988	N	LEU	A	70	34.923	87.802	-5.012	1.00	29.22	T4
	ATOM	3989	CA	LEU	A	70	35.396	89.030	-5.627	1.00	29.57	T4
	ATOM	3990	CB	LEU	A	70	36.874	88.912	-5.969	1.00	39.75	T4
	ATOM	3991	CG	LEU	A	70	37.799	88.219	-4.982	1.00	29.80	T4
50	ATOM	3992	CD1	LEU	A	70	39.147	88.041	-5.632	1.00	34.22	T4
	ATOM	3993	CD2	LEU	A	70	37.908	89.032	-3.709	1.00	36.37	T4
	ATOM	3994	C	LEU	A	70	34.664	89.298	-6.926	1.00	33.06	T4
	ATOM	3995	O	LEU	A	70	34.332	88.370	-7.657	1.00	32.94	T4
	ATOM	3996	N	ILE	A	71	34.410	90.568	-7.213	1.00	25.58	T4
55	ATOM	3997	CA	ILE	A	71	33.782	90.928	-8.469	1.00	33.51	T4
	ATOM	3998	CB	ILE	A	71	32.603	91.886	-8.263	1.00	35.54	T4
	ATOM	3999	CG2	ILE	A	71	32.171	92.475	-9.590	1.00	22.16	T4
	ATOM	4000	CG1	ILE	A	71	31.432	91.128	-7.645	1.00	24.83	T4
	ATOM	4001	CD1	ILE	A	71	30.222	91.988	-7.366	1.00	30.81	T4
60	ATOM	4002	C	ILE	A	71	34.915	91.615	-9.217	1.00	22.72	T4
	ATOM	4003	O	ILE	A	71	35.322	92.710	-8.861	1.00	31.50	T4
	ATOM	4004	N	GLN	A	72	35.430	90.971	-10.255	1.00	33.82	T4
	ATOM	4005	CA	GLN	A	72	36.549	91.533	-10.989	1.00	36.51	T4
	ATOM	4006	CB	GLN	A	72	37.667	90.509	-11.038	1.00	29.80	T4
65	ATOM	4007	CG	GLN	A	72	37.902	89.872	-9.694	1.00	30.42	T4
	ATOM	4008	CD	GLN	A	72	39.134	88.995	-9.673	1.00	29.07	T4

	ATOM	4009	OE1	GLN	A	72	39.285	88.091	-10.502	1.00	30.99	T4
	ATOM	4010	NE2	GLN	A	72	40.031	89.257	-8.719	1.00	24.65	T4
	ATOM	4011	C	GLN	A	72	36.273	92.039	-12.389	1.00	28.98	T4
	ATOM	4012	O	GLN	A	72	35.328	91.623	-13.051	1.00	26.96	T4
5	ATOM	4013	N	ARG	A	73	37.143	92.938	-12.831	1.00	32.82	T4
	ATOM	4014	CA	ARG	A	73	37.046	93.547	-14.143	1.00	31.73	T4
	ATOM	4015	CB	ARG	A	73	36.836	95.046	-13.971	1.00	26.99	T4
	ATOM	4016	CG	ARG	A	73	36.745	95.790	-15.266	1.00	32.59	T4
	ATOM	4017	CD	ARG	A	73	37.135	97.226	-15.084	1.00	24.66	T4
10	ATOM	4018	NE	ARG	A	73	37.067	97.951	-16.347	1.00	32.91	T4
	ATOM	4019	CZ	ARG	A	73	37.600	99.149	-16.550	1.00	32.58	T4
	ATOM	4020	NH1	ARG	A	73	38.251	99.762	-15.570	1.00	24.49	T4
	ATOM	4021	NH2	ARG	A	73	37.474	99.734	-17.729	1.00	33.02	T4
	ATOM	4022	C	ARG	A	73	38.312	93.302	-14.971	1.00	29.68	T4
15	ATOM	4023	O	ARG	A	73	39.421	93.486	-14.473	1.00	34.62	T4
	ATOM	4024	N	LYS	A	74	38.146	92.882	-16.225	1.00	27.35	T4
	ATOM	4025	CA	LYS	A	74	39.281	92.655	-17.118	1.00	31.91	T4
	ATOM	4026	CB	LYS	A	74	39.147	91.330	-17.861	1.00	25.38	T4
	ATOM	4027	CG	LYS	A	74	39.275	90.102	-16.980	1.00	26.92	T4
20	ATOM	4028	CD	LYS	A	74	39.023	88.792	-17.757	1.00	29.18	T4
	ATOM	4029	CE	LYS	A	74	39.120	87.560	-16.831	1.00	27.26	T4
	ATOM	4030	NZ	LYS	A	74	38.943	86.259	-17.560	1.00	34.38	T4
	ATOM	4031	C	LYS	A	74	39.322	93.783	-18.135	1.00	29.17	T4
	ATOM	4032	O	LYS	A	74	38.569	93.771	-19.105	1.00	26.33	T4
25	ATOM	4033	N	LYS	A	75	40.196	94.757	-17.910	1.00	35.07	T4
	ATOM	4034	CA	LYS	A	75	40.330	95.894	-18.813	1.00	29.95	T4
	ATOM	4035	CB	LYS	A	75	41.411	96.846	-18.307	1.00	35.76	T4
	ATOM	4036	CG	LYS	A	75	41.178	97.436	-16.932	1.00	23.77	T4
	ATOM	4037	CD	LYS	A	75	42.334	98.371	-16.569	1.00	30.09	T4
30	ATOM	4038	CE	LYS	A	75	42.147	98.976	-15.172	1.00	29.52	T4
	ATOM	4039	NZ	LYS	A	75	43.306	99.810	-14.691	1.00	31.16	T4
	ATOM	4040	C	LYS	A	75	40.707	95.467	-20.227	1.00	34.93	T4
	ATOM	4041	O	LYS	A	75	41.574	94.621	-20.413	1.00	33.31	T4
	ATOM	4042	N	VAL	A	76	40.071	96.065	-21.226	1.00	33.96	T4
35	ATOM	4043	CA	VAL	A	76	40.379	95.742	-22.618	1.00	25.57	T4
	ATOM	4044	CB	VAL	A	76	39.330	96.255	-23.596	1.00	41.06	T4
	ATOM	4045	CG1	VAL	A	76	39.487	95.562	-24.911	1.00	29.96	T4
	ATOM	4046	CG2	VAL	A	76	37.966	96.057	-23.049	1.00	29.38	T4
	ATOM	4047	C	VAL	A	76	41.641	96.494	-22.973	1.00	33.64	T4
40	ATOM	4048	O	VAL	A	76	42.523	95.978	-23.661	1.00	35.69	T4
	ATOM	4049	N	HIS	A	77	41.699	97.737	-22.511	1.00	33.53	T4
	ATOM	4050	CA	HIS	A	77	42.834	98.600	-22.758	1.00	27.78	T4
	ATOM	4051	CB	HIS	A	77	42.341	99.973	-23.172	1.00	32.26	T4
	ATOM	4052	CG	HIS	A	77	41.477	99.950	-24.391	1.00	28.33	T4
45	ATOM	4053	CD2	HIS	A	77	40.576	100.844	-24.864	1.00	31.69	T4
	ATOM	4054	ND1	HIS	A	77	41.542	98.938	-25.327	1.00	29.99	T4
	ATOM	4055	CE1	HIS	A	77	40.722	99.210	-26.326	1.00	34.74	T4
	ATOM	4056	NE2	HIS	A	77	40.125	100.361	-26.071	1.00	33.71	T4
	ATOM	4057	C	HIS	A	77	43.672	98.689	-21.494	1.00	23.34	T4
50	ATOM	4058	O	HIS	A	77	43.127	98.658	-20.384	1.00	30.97	T4
	ATOM	4059	N	VAL	A	78	44.992	98.816	-21.657	1.00	33.18	T4
	ATOM	4060	CA	VAL	A	78	45.856	98.854	-20.497	1.00	28.93	T4
	ATOM	4061	CB	VAL	A	78	46.652	97.537	-20.420	1.00	23.28	T4
	ATOM	4062	CG1	VAL	A	78	47.430	97.468	-19.115	1.00	26.51	T4
55	ATOM	4063	CG2	VAL	A	78	45.689	96.367	-20.468	1.00	31.54	T4
	ATOM	4064	C	VAL	A	78	46.779	100.056	-20.236	1.00	29.98	T4
	ATOM	4065	O	VAL	A	78	46.574	100.775	-19.250	1.00	33.90	T4
	ATOM	4066	N	PHE	A	79	47.786	100.296	-21.066	1.00	29.24	T4
	ATOM	4067	CA	PHE	A	79	48.700	101.432	-20.813	1.00	36.25	T4
60	ATOM	4068	CB	PHE	A	79	47.944	102.711	-20.415	1.00	30.06	T4
	ATOM	4069	CG	PHE	A	79	46.852	103.100	-21.353	1.00	25.74	T4
	ATOM	4070	CD1	PHE	A	79	45.526	102.822	-21.051	1.00	38.38	T4
	ATOM	4071	CD2	PHE	A	79	47.145	103.747	-22.543	1.00	36.48	T4
	ATOM	4072	CE1	PHE	A	79	44.511	103.178	-21.912	1.00	40.36	T4
65	ATOM	4073	CE2	PHE	A	79	46.137	104.110	-23.420	1.00	27.70	T4
	ATOM	4074	CZ	PHE	A	79	44.816	103.824	-23.102	1.00	32.14	T4

	ATOM	4075	C	PHE	A	79	49.741	101.198	-19.706	1.00	38.96	T4
	ATOM	4076	O	PHE	A	79	49.396	100.919	-18.554	1.00	32.50	T4
	ATOM	4077	N	GLY	A	80	51.014	101.355	-20.063	1.00	32.15	T4
	ATOM	4078	CA	GLY	A	80	52.102	101.206	-19.109	1.00	32.64	T4
5	ATOM	4079	C	GLY	A	80	52.063	99.993	-18.206	1.00	28.84	T4
	ATOM	4080	O	GLY	A	80	51.878	98.864	-18.665	1.00	38.31	T4
	ATOM	4081	N	ASP	A	81	52.231	100.235	-16.909	1.00	33.18	T4
	ATOM	4082	CA	ASP	A	81	52.246	99.170	-15.920	1.00	32.36	T4
	ATOM	4083	CB	ASP	A	81	53.362	99.430	-14.903	1.00	23.34	T4
10	ATOM	4084	CG	ASP	A	81	53.123	100.667	-14.065	1.00	35.88	T4
	ATOM	4085	OD1	ASP	A	81	52.287	101.499	-14.454	1.00	29.26	T4
	ATOM	4086	OD2	ASP	A	81	53.786	100.815	-13.017	1.00	29.04	T4
	ATOM	4087	C	ASP	A	81	50.926	98.945	-15.197	1.00	30.18	T4
	ATOM	4088	O	ASP	A	81	50.908	98.521	-14.045	1.00	33.90	T4
15	ATOM	4089	N	GLU	A	82	49.817	99.230	-15.865	1.00	29.28	T4
	ATOM	4090	CA	GLU	A	82	48.513	99.018	-15.256	1.00	27.59	T4
	ATOM	4091	CB	GLU	A	82	47.398	99.574	-16.131	1.00	34.71	T4
	ATOM	4092	CG	GLU	A	82	47.150	101.045	-16.076	1.00	28.61	T4
	ATOM	4093	CD	GLU	A	82	45.700	101.346	-16.394	1.00	29.08	T4
20	ATOM	4094	OE1	GLU	A	82	45.177	100.744	-17.349	1.00	33.65	T4
	ATOM	4095	OE2	GLU	A	82	45.075	102.172	-15.690	1.00	37.95	T4
	ATOM	4096	C	GLU	A	82	48.287	97.526	-15.185	1.00	30.95	T4
	ATOM	4097	O	GLU	A	82	48.788	96.800	-16.034	1.00	24.75	T4
	ATOM	4098	N	LEU	A	83	47.542	97.058	-14.191	1.00	29.74	T4
25	ATOM	4099	CA	LEU	A	83	47.230	95.639	-14.127	1.00	28.37	T4
	ATOM	4100	CB	LEU	A	83	47.045	95.159	-12.692	1.00	27.58	T4
	ATOM	4101	CG	LEU	A	83	48.285	95.107	-11.816	1.00	25.75	T4
	ATOM	4102	CD1	LEU	A	83	48.817	96.512	-11.617	1.00	29.25	T4
	ATOM	4103	CD2	LEU	A	83	47.930	94.488	-10.487	1.00	30.41	T4
30	ATOM	4104	C	LEU	A	83	45.904	95.593	-14.857	1.00	26.98	T4
	ATOM	4105	O	LEU	A	83	45.012	96.397	-14.577	1.00	31.30	T4
	ATOM	4106	N	SER	A	84	45.773	94.675	-15.806	1.00	28.79	T4
	ATOM	4107	CA	SER	A	84	44.541	94.580	-16.582	1.00	30.90	T4
	ATOM	4108	CB	SER	A	84	44.804	93.865	-17.916	1.00	27.45	T4
35	ATOM	4109	OG	SER	A	84	45.558	92.682	-17.733	1.00	28.80	T4
	ATOM	4110	C	SER	A	84	43.408	93.900	-15.835	1.00	28.39	T4
	ATOM	4111	O	SER	A	84	42.287	93.839	-16.333	1.00	35.53	T4
	ATOM	4112	N	LEU	A	85	43.698	93.403	-14.637	1.00	28.38	T4
	ATOM	4113	CA	LEU	A	85	42.685	92.741	-13.832	1.00	31.64	T4
40	ATOM	4114	CB	LEU	A	85	43.111	91.302	-13.525	1.00	31.76	T4
	ATOM	4115	CG	LEU	A	85	42.086	90.273	-13.026	1.00	33.80	T4
	ATOM	4116	CD1	LEU	A	85	41.635	90.586	-11.612	1.00	35.60	T4
	ATOM	4117	CD2	LEU	A	85	40.909	90.241	-13.974	1.00	28.60	T4
	ATOM	4118	C	LEU	A	85	42.496	93.536	-12.547	1.00	32.40	T4
45	ATOM	4119	O	LEU	A	85	43.352	93.536	-11.670	1.00	28.53	T4
	ATOM	4120	N	VAL	A	86	41.375	94.241	-12.454	1.00	33.55	T4
	ATOM	4121	CA	VAL	A	86	41.065	95.035	-11.266	1.00	30.15	T4
	ATOM	4122	CB	VAL	A	86	40.603	96.461	-11.623	1.00	32.17	T4
	ATOM	4123	CG1	VAL	A	86	40.206	97.203	-10.361	1.00	36.77	T4
50	ATOM	4124	CG2	VAL	A	86	41.707	97.202	-12.355	1.00	28.26	T4
	ATOM	4125	C	VAL	A	86	39.924	94.367	-10.519	1.00	32.09	T4
	ATOM	4126	O	VAL	A	86	38.968	93.885	-11.125	1.00	32.24	T4
	ATOM	4127	N	THR	A	87	40.024	94.327	-9.202	1.00	30.20	T4
	ATOM	4128	CA	THR	A	87	38.966	93.726	-8.439	1.00	37.28	T4
55	ATOM	4129	CB	THR	A	87	39.505	92.679	-7.425	1.00	27.50	T4
	ATOM	4130	OG1	THR	A	87	39.221	93.104	-6.092	1.00	30.85	T4
	ATOM	4131	CG2	THR	A	87	40.994	92.479	-7.596	1.00	23.62	T4
	ATOM	4132	C	THR	A	87	38.211	94.835	-7.734	1.00	38.78	T4
	ATOM	4133	O	THR	A	87	38.772	95.579	-6.944	1.00	29.55	T4
60	ATOM	4134	N	LEU	A	88	36.933	94.956	-8.072	1.00	31.34	T4
	ATOM	4135	CA	LEU	A	88	36.037	95.948	-7.496	1.00	22.58	T4
	ATOM	4136	CB	LEU	A	88	35.031	96.405	-8.559	1.00	27.22	T4
	ATOM	4137	CG	LEU	A	88	35.394	96.981	-9.935	1.00	34.39	T4
	ATOM	4138	CD1	LEU	A	88	36.684	96.441	-10.424	1.00	26.61	T4
65	ATOM	4139	CD2	LEU	A	88	34.312	96.632	-10.914	1.00	36.79	T4
	ATOM	4140	C	LEU	A	88	35.251	95.257	-6.378	1.00	31.93	T4

	ATOM	4141	O	LEU	A	88	34.939	94.066	-6.456	1.00	28.01	T4
	ATOM	4142	N	PHE	A	89	34.941	95.969	-5.314	1.00	27.23	T4
	ATOM	4143	CA	PHE	A	89	34.098	95.355	-4.279	1.00	31.67	T4
	ATOM	4144	CB	PHE	A	89	32.771	94.935	-4.921	1.00	27.26	T4
5	ATOM	4145	CG	PHE	A	89	32.233	95.970	-5.874	1.00	36.28	T4
	ATOM	4146	CD1	PHE	A	89	31.552	95.595	-7.024	1.00	32.61	T4
	ATOM	4147	CD2	PHE	A	89	32.509	97.332	-5.663	1.00	31.95	T4
	ATOM	4148	CE1	PHE	A	89	31.165	96.557	-7.954	1.00	28.86	T4
	ATOM	4149	CE2	PHE	A	89	32.126	98.300	-6.586	1.00	34.94	T4
10	ATOM	4150	CZ	PHE	A	89	31.456	97.914	-7.735	1.00	28.87	T4
	ATOM	4151	C	PHE	A	89	34.645	94.216	-3.429	1.00	33.39	T4
	ATOM	4152	O	PHE	A	89	35.430	94.473	-2.520	1.00	23.89	T4
	ATOM	4153	N	ARG	A	90	34.223	92.976	-3.666	1.00	33.10	T4
	ATOM	4154	CA	ARG	A	90	34.713	91.874	-2.816	1.00	26.81	T4
15	ATOM	4155	CB	ARG	A	90	36.244	91.860	-2.811	1.00	27.47	T4
	ATOM	4156	CG	ARG	A	90	36.864	91.280	-1.543	1.00	28.81	T4
	ATOM	4157	CD	ARG	A	90	38.315	91.663	-1.426	1.00	37.30	T4
	ATOM	4158	NE	ARG	A	90	38.835	91.284	-0.125	1.00	34.40	T4
	ATOM	4159	CZ	ARG	A	90	39.719	91.998	0.565	1.00	27.86	T4
20	ATOM	4160	NH1	ARG	A	90	40.193	93.144	0.072	1.00	28.06	T4
	ATOM	4161	NH2	ARG	A	90	40.109	91.576	1.766	1.00	30.88	T4
	ATOM	4162	C	ARG	A	90	34.231	91.920	-1.342	1.00	27.09	T4
	ATOM	4163	O	ARG	A	90	34.378	92.931	-0.658	1.00	33.18	T4
	ATOM	4164	N	CYS	A	91	33.689	90.805	-0.856	1.00	27.62	T4
25	ATOM	4165	CA	CYS	A	91	33.191	90.711	0.519	1.00	38.34	T4
	ATOM	4166	CB	CYS	A	91	31.665	90.688	0.525	1.00	27.74	T4
	ATOM	4167	SG	CYS	A	91	30.965	89.371	-0.475	1.00	32.02	T4
	ATOM	4168	C	CYS	A	91	33.725	89.477	1.250	1.00	33.87	T4
	ATOM	4169	O	CYS	A	91	34.264	88.566	0.630	1.00	31.62	T4
30	ATOM	4170	N	ILE	A	92	33.562	89.450	2.572	1.00	34.41	T4
	ATOM	4171	CA	ILE	A	92	34.065	88.349	3.402	1.00	35.08	T4
	ATOM	4172	CB	ILE	A	92	35.368	88.747	4.126	1.00	28.49	T4
	ATOM	4173	CG2	ILE	A	92	35.908	87.585	4.908	1.00	32.23	T4
	ATOM	4174	CG1	ILE	A	92	36.411	89.202	3.114	1.00	26.15	T4
35	ATOM	4175	CD1	ILE	A	92	37.599	89.905	3.742	1.00	33.02	T4
	ATOM	4176	C	ILE	A	92	33.059	88.006	4.478	1.00	29.79	T4
	ATOM	4177	O	ILE	A	92	32.280	88.853	4.873	1.00	28.60	T4
	ATOM	4178	N	GLN	A	93	33.084	86.768	4.958	1.00	36.61	T4
	ATOM	4179	CA	GLN	A	93	32.174	86.334	6.022	1.00	27.08	T4
40	ATOM	4180	CB	GLN	A	93	30.847	85.842	5.432	1.00	25.15	T4
	ATOM	4181	CG	GLN	A	93	29.813	86.933	5.189	1.00	28.09	T4
	ATOM	4182	CD	GLN	A	93	28.803	87.054	6.325	1.00	27.82	T4
	ATOM	4183	OE1	GLN	A	93	29.095	87.575	7.401	1.00	30.63	T4
	ATOM	4184	NE2	GLN	A	93	27.598	86.555	6.082	1.00	32.61	T4
45	ATOM	4185	C	GLN	A	93	32.796	85.228	6.872	1.00	34.32	T4
	ATOM	4186	O	GLN	A	93	33.172	84.168	6.359	1.00	35.08	T4
	ATOM	4187	N	ASN	A	94	32.924	85.480	8.169	1.00	23.60	T4
	ATOM	4188	CA	ASN	A	94	33.488	84.473	9.050	1.00	33.48	T4
	ATOM	4189	CB	ASN	A	94	33.592	85.003	10.486	1.00	23.43	T4
50	ATOM	4190	CG	ASN	A	94	34.828	85.869	10.705	1.00	31.05	T4
	ATOM	4191	OD1	ASN	A	94	35.951	85.441	10.433	1.00	29.40	T4
	ATOM	4192	ND2	ASN	A	94	34.627	87.085	11.204	1.00	33.22	T4
	ATOM	4193	C	ASN	A	94	32.541	83.289	8.995	1.00	35.59	T4
	ATOM	4194	O	ASN	A	94	31.340	83.467	8.844	1.00	24.81	T4
55	ATOM	4195	N	MET	A	95	33.077	82.082	9.097	1.00	38.34	T4
	ATOM	4196	CA	MET	A	95	32.253	80.883	9.054	1.00	24.52	T4
	ATOM	4197	CB	MET	A	95	32.738	79.938	7.948	1.00	24.06	T4
	ATOM	4198	CG	MET	A	95	32.676	80.518	6.547	1.00	28.84	T4
	ATOM	4199	SD	MET	A	95	31.041	81.174	6.151	1.00	36.47	T4
60	ATOM	4200	CE	MET	A	95	30.071	79.686	5.898	1.00	28.85	T4
	ATOM	4201	C	MET	A	95	32.296	80.146	10.386	1.00	30.86	T4
	ATOM	4202	O	MET	A	95	33.284	80.224	11.115	1.00	32.55	T4
	ATOM	4203	N	PRO	A	96	31.213	79.424	10.721	1.00	31.95	T4
	ATOM	4204	CD	PRO	A	96	29.929	79.364	10.010	1.00	29.48	T4
65	ATOM	4205	CA	PRO	A	96	31.127	78.663	11.967	1.00	31.04	T4
	ATOM	4206	CB	PRO	A	96	29.635	78.345	12.100	1.00	31.15	T4

	ATOM	4207	CG	PRO	A	96	28.967	79.290	11.162	1.00	31.30	T4
	ATOM	4208	C	PRO	A	96	31.923	77.398	11.712	1.00	28.64	T4
	ATOM	4209	O	PRO	A	96	32.614	77.282	10.692	1.00	24.52	T4
	ATOM	4210	N	GLU	A	97	31.805	76.436	12.617	1.00	32.86	T4
5	ATOM	4211	CA	GLU	A	97	32.530	75.196	12.451	1.00	37.32	T4
	ATOM	4212	CB	GLU	A	97	33.379	74.946	13.682	1.00	29.84	T4
	ATOM	4213	CG	GLU	A	97	34.441	73.897	13.463	1.00	30.29	T4
	ATOM	4214	CD	GLU	A	97	35.784	74.328	14.046	1.00	32.94	T4
	ATOM	4215	OE1	GLU	A	97	35.847	74.544	15.284	1.00	28.86	T4
10	ATOM	4216	OE2	GLU	A	97	36.773	74.463	13.270	1.00	24.92	T4
	ATOM	4217	C	GLU	A	97	31.561	74.048	12.250	1.00	26.65	T4
	ATOM	4218	O	GLU	A	97	31.902	73.014	11.676	1.00	32.39	T4
	ATOM	4219	N	THR	A	98	30.336	74.254	12.710	1.00	29.20	T4
	ATOM	4220	CA	THR	A	98	29.321	73.226	12.632	1.00	27.66	T4
15	ATOM	4221	CB	THR	A	98	28.436	73.268	13.870	1.00	27.12	T4
	ATOM	4222	OG1	THR	A	98	27.756	74.532	13.912	1.00	24.94	T4
	ATOM	4223	CG2	THR	A	98	29.286	73.112	15.129	1.00	37.45	T4
	ATOM	4224	C	THR	A	98	28.424	73.272	11.405	1.00	31.43	T4
	ATOM	4225	O	THR	A	98	28.560	72.433	10.514	1.00	32.28	T4
20	ATOM	4226	N	LEU	A	99	27.512	74.236	11.336	1.00	33.69	T4
	ATOM	4227	CA	LEU	A	99	26.606	74.260	10.200	1.00	32.18	T4
	ATOM	4228	CB	LEU	A	99	25.175	74.114	10.708	1.00	34.54	T4
	ATOM	4229	CG	LEU	A	99	24.946	72.777	11.421	1.00	29.76	T4
	ATOM	4230	CD1	LEU	A	99	23.566	72.722	12.046	1.00	34.05	T4
25	ATOM	4231	CD2	LEU	A	99	25.119	71.658	10.421	1.00	27.41	T4
	ATOM	4232	C	LEU	A	99	26.736	75.477	9.302	1.00	31.61	T4
	ATOM	4233	O	LEU	A	99	25.863	76.353	9.283	1.00	25.51	T4
	ATOM	4234	N	PRO	A	100	27.825	75.539	8.518	1.00	34.43	T4
	ATOM	4235	CD	PRO	A	100	28.868	74.510	8.404	1.00	29.15	T4
30	ATOM	4236	CA	PRO	A	100	28.101	76.644	7.599	1.00	30.77	T4
	ATOM	4237	CB	PRO	A	100	29.228	76.093	6.737	1.00	25.65	T4
	ATOM	4238	CG	PRO	A	100	29.982	75.269	7.712	1.00	23.26	T4
	ATOM	4239	C	PRO	A	100	26.894	77.058	6.774	1.00	33.67	T4
	ATOM	4240	O	PRO	A	100	26.315	76.245	6.056	1.00	27.12	T4
35	ATOM	4241	N	ASN	A	101	26.528	78.329	6.889	1.00	31.73	T4
	ATOM	4242	CA	ASN	A	101	25.400	78.877	6.163	1.00	30.24	T4
	ATOM	4243	CB	ASN	A	101	24.098	78.446	6.803	1.00	24.83	T4
	ATOM	4244	CG	ASN	A	101	23.618	77.143	6.279	1.00	31.71	T4
	ATOM	4245	OD1	ASN	A	101	23.217	77.045	5.117	1.00	30.18	T4
40	ATOM	4246	ND2	ASN	A	101	23.663	76.111	7.121	1.00	28.83	T4
	ATOM	4247	C	ASN	A	101	25.454	80.372	6.205	1.00	39.00	T4
	ATOM	4248	O	ASN	A	101	24.780	80.974	7.042	1.00	30.76	T4
	ATOM	4249	N	ASN	A	102	26.225	80.997	5.318	1.00	26.75	T4
	ATOM	4250	CA	ASN	A	102	26.262	82.439	5.394	1.00	34.72	T4
45	ATOM	4251	CB	ASN	A	102	27.633	82.907	5.867	1.00	32.28	T4
	ATOM	4252	CG	ASN	A	102	27.742	82.876	7.390	1.00	31.70	T4
	ATOM	4253	OD1	ASN	A	102	26.831	83.330	8.096	1.00	30.02	T4
	ATOM	4254	ND2	ASN	A	102	28.848	82.333	7.901	1.00	29.80	T4
	ATOM	4255	C	ASN	A	102	25.745	83.316	4.277	1.00	32.06	T4
50	ATOM	4256	O	ASN	A	102	24.988	84.241	4.564	1.00	33.48	T4
	ATOM	4257	N	SER	A	103	26.093	83.082	3.022	1.00	32.64	T4
	ATOM	4258	CA	SER	A	103	25.533	83.986	2.009	1.00	31.99	T4
	ATOM	4259	CB	SER	A	103	23.986	83.908	2.020	1.00	33.64	T4
	ATOM	4260	OG	SER	A	103	23.372	85.195	2.020	1.00	32.11	T4
55	ATOM	4261	C	SER	A	103	25.976	85.446	2.245	1.00	31.18	T4
	ATOM	4262	O	SER	A	103	25.853	85.999	3.344	1.00	35.69	T4
	ATOM	4263	N	CYS	A	104	26.477	86.082	1.200	1.00	29.04	T4
	ATOM	4264	CA	CYS	A	104	26.933	87.442	1.349	1.00	32.22	T4
	ATOM	4265	CB	CYS	A	104	28.415	87.429	1.702	1.00	23.20	T4
60	ATOM	4266	SG	CYS	A	104	29.087	88.990	2.204	1.00	32.89	T4
	ATOM	4267	C	CYS	A	104	26.694	88.205	0.064	1.00	30.43	T4
	ATOM	4268	O	CYS	A	104	27.081	87.765	-1.011	1.00	36.87	T4
	ATOM	4269	N	TYR	A	105	26.039	89.349	0.184	1.00	26.72	T4
	ATOM	4270	CA	TYR	A	105	25.732	90.193	-0.960	1.00	31.43	T4
65	ATOM	4271	CB	TYR	A	105	24.265	90.627	-0.906	1.00	32.49	T4
	ATOM	4272	CG	TYR	A	105	23.847	91.600	-1.985	1.00	30.12	T4

	ATOM	4273	CD1	TYR	A	105	23.283	91.150	-3.170	1.00	32.66	T4
	ATOM	4274	CE1	TYR	A	105	22.888	92.043	-4.162	1.00	28.21	T4
	ATOM	4275	CD2	TYR	A	105	24.011	92.973	-1.818	1.00	37.83	T4
	ATOM	4276	CE2	TYR	A	105	23.624	93.872	-2.802	1.00	30.33	T4
5	ATOM	4277	CZ	TYR	A	105	23.063	93.401	-3.966	1.00	30.69	T4
	ATOM	4278	OH	TYR	A	105	22.661	94.288	-4.930	1.00	31.57	T4
	ATOM	4279	C	TYR	A	105	26.614	91.426	-0.925	1.00	28.55	T4
	ATOM	4280	O	TYR	A	105	26.953	91.924	0.143	1.00	29.48	T4
	ATOM	4281	N	SER	A	106	26.984	91.922	-2.096	1.00	32.17	T4
10	ATOM	4282	CA	SER	A	106	27.800	93.124	-2.180	1.00	35.86	T4
	ATOM	4283	CB	SER	A	106	29.274	92.800	-1.913	1.00	25.82	T4
	ATOM	4284	OG	SER	A	106	30.048	93.972	-1.754	1.00	28.96	T4
	ATOM	4285	C	SER	A	106	27.610	93.676	-3.580	1.00	29.39	T4
	ATOM	4286	O	SER	A	106	27.464	92.913	-4.533	1.00	33.12	T4
15	ATOM	4287	N	ALA	A	107	27.591	94.999	-3.696	1.00	28.21	T4
	ATOM	4288	CA	ALA	A	107	27.405	95.649	-4.982	1.00	31.24	T4
	ATOM	4289	CB	ALA	A	107	25.925	95.720	-5.310	1.00	35.90	T4
	ATOM	4290	C	ALA	A	107	28.010	97.049	-4.987	1.00	27.29	T4
	ATOM	4291	O	ALA	A	107	28.324	97.605	-3.940	1.00	30.12	T4
20	ATOM	4292	N	GLY	A	108	28.170	97.611	-6.179	1.00	33.53	T4
	ATOM	4293	CA	GLY	A	108	28.727	98.942	-6.309	1.00	33.83	T4
	ATOM	4294	C	GLY	A	108	28.640	99.441	-7.737	1.00	34.81	T4
	ATOM	4295	O	GLY	A	108	28.164	98.733	-8.614	1.00	27.66	T4
	ATOM	4296	N	ILE	A	109	29.096	100.664	-7.972	1.00	25.89	T4
25	ATOM	4297	CA	ILE	A	109	29.071	101.248	-9.306	1.00	34.11	T4
	ATOM	4298	CB	ILE	A	109	28.432	102.642	-9.287	1.00	28.90	T4
	ATOM	4299	CG2	ILE	A	109	28.453	103.237	-10.677	1.00	33.85	T4
	ATOM	4300	CG1	ILE	A	109	27.003	102.553	-8.765	1.00	27.45	T4
	ATOM	4301	CD1	ILE	A	109	26.372	103.888	-8.519	1.00	31.51	T4
30	ATOM	4302	C	ILE	A	109	30.495	101.384	-9.831	1.00	32.93	T4
	ATOM	4303	O	ILE	A	109	31.413	101.710	-9.081	1.00	31.22	T4
	ATOM	4304	N	ALA	A	110	30.680	101.136	-11.119	1.00	33.61	T4
	ATOM	4305	CA	ALA	A	110	31.998	101.234	-11.723	1.00	30.68	T4
	ATOM	4306	CB	ALA	A	110	32.704	99.899	-11.643	1.00	27.21	T4
35	ATOM	4307	C	ALA	A	110	31.858	101.642	-13.173	1.00	32.48	T4
	ATOM	4308	O	ALA	A	110	30.820	101.406	-13.787	1.00	32.78	T4
	ATOM	4309	N	LYS	A	111	32.893	102.264	-13.723	1.00	26.76	T4
	ATOM	4310	CA	LYS	A	111	32.836	102.659	-15.115	1.00	28.46	T4
	ATOM	4311	CB	LYS	A	111	33.554	103.979	-15.348	1.00	29.12	T4
40	ATOM	4312	CG	LYS	A	111	33.452	104.425	-16.805	1.00	27.14	T4
	ATOM	4313	CD	LYS	A	111	34.119	105.766	-17.037	1.00	23.18	T4
	ATOM	4314	CE	LYS	A	111	33.966	106.218	-18.494	1.00	30.17	T4
	ATOM	4315	NZ	LYS	A	111	34.593	107.573	-18.731	1.00	29.95	T4
	ATOM	4316	C	LYS	A	111	33.508	101.561	-15.914	1.00	26.86	T4
45	ATOM	4317	O	LYS	A	111	34.589	101.118	-15.550	1.00	35.06	T4
	ATOM	4318	N	LEU	A	112	32.862	101.125	-16.995	1.00	32.48	T4
	ATOM	4319	CA	LEU	A	112	33.379	100.051	-17.833	1.00	32.94	T4
	ATOM	4320	CB	LEU	A	112	32.496	98.812	-17.681	1.00	33.30	T4
	ATOM	4321	CG	LEU	A	112	32.176	98.355	-16.263	1.00	32.88	T4
50	ATOM	4322	CD1	LEU	A	112	31.070	97.326	-16.294	1.00	21.72	T4
	ATOM	4323	CD2	LEU	A	112	33.419	97.802	-15.620	1.00	30.45	T4
	ATOM	4324	C	LEU	A	112	33.394	100.471	-19.295	1.00	30.81	T4
	ATOM	4325	O	LEU	A	112	32.717	101.417	-19.666	1.00	29.74	T4
	ATOM	4326	N	GLU	A	113	34.161	99.753	-20.118	1.00	25.19	T4
55	ATOM	4327	CA	GLU	A	113	34.256	100.044	-21.548	1.00	25.11	T4
	ATOM	4328	CB	GLU	A	113	35.635	100.562	-21.924	1.00	34.00	T4
	ATOM	4329	CG	GLU	A	113	36.281	101.497	-20.961	1.00	31.90	T4
	ATOM	4330	CD	GLU	A	113	37.392	102.291	-21.629	1.00	36.72	T4
	ATOM	4331	OE1	GLU	A	113	38.203	101.688	-22.396	1.00	29.76	T4
60	ATOM	4332	OE2	GLU	A	113	37.451	103.528	-21.382	1.00	30.99	T4
	ATOM	4333	C	GLU	A	113	34.016	98.821	-22.413	1.00	29.21	T4
	ATOM	4334	O	GLU	A	113	34.197	97.688	-21.970	1.00	27.13	T4
	ATOM	4335	N	GLU	A	114	33.635	99.069	-23.662	1.00	33.13	T4
	ATOM	4336	CA	GLU	A	114	33.411	98.002	-24.625	1.00	31.60	T4
65	ATOM	4337	CB	GLU	A	114	33.445	98.534	-26.044	1.00	32.36	T4
	ATOM	4338	CG	GLU	A	114	32.159	99.015	-26.606	1.00	29.76	T4

	ATOM	4339	CD	GLU	A	114	32.233	99.068	-28.115	1.00	32.78	T4
	ATOM	4340	OE1	GLU	A	114	32.389	97.991	-28.732	1.00	33.16	T4
	ATOM	4341	OE2	GLU	A	114	32.149	100.179	-28.678	1.00	31.35	T4
	ATOM	4342	C	GLU	A	114	34.542	96.998	-24.535	1.00	29.28	T4
5	ATOM	4343	O	GLU	A	114	35.708	97.366	-24.668	1.00	27.75	T4
	ATOM	4344	N	GLY	A	115	34.208	95.732	-24.336	1.00	27.03	T4
	ATOM	4345	CA	GLY	A	115	35.244	94.721	-24.277	1.00	35.74	T4
	ATOM	4346	C	GLY	A	115	35.635	94.306	-22.882	1.00	33.81	T4
	ATOM	4347	O	GLY	A	115	36.257	93.261	-22.708	1.00	31.56	T4
10	ATOM	4348	N	ASP	A	116	35.309	95.117	-21.884	1.00	37.93	T4
	ATOM	4349	CA	ASP	A	116	35.655	94.737	-20.531	1.00	24.66	T4
	ATOM	4350	CB	ASP	A	116	35.331	95.860	-19.540	1.00	27.42	T4
	ATOM	4351	CG	ASP	A	116	36.278	97.044	-19.647	1.00	38.04	T4
	ATOM	4352	OD1	ASP	A	116	37.427	96.859	-20.086	1.00	32.47	T4
15	ATOM	4353	OD2	ASP	A	116	35.882	98.163	-19.264	1.00	35.93	T4
	ATOM	4354	C	ASP	A	116	34.831	93.507	-20.186	1.00	32.30	T4
	ATOM	4355	O	ASP	A	116	33.758	93.292	-20.755	1.00	32.72	T4
	ATOM	4356	N	GLU	A	117	35.333	92.690	-19.271	1.00	37.37	T4
	ATOM	4357	CA	GLU	A	117	34.592	91.515	-18.856	1.00	26.93	T4
20	ATOM	4358	CB	GLU	A	117	35.295	90.240	-19.317	1.00	31.18	T4
	ATOM	4359	CG	GLU	A	117	35.570	90.190	-20.798	1.00	29.49	T4
	ATOM	4360	CD	GLU	A	117	36.189	88.868	-21.222	1.00	28.88	T4
	ATOM	4361	OE1	GLU	A	117	37.005	88.309	-20.443	1.00	37.05	T4
	ATOM	4362	OE2	GLU	A	117	35.866	88.396	-22.342	1.00	20.40	T4
25	ATOM	4363	C	GLU	A	117	34.494	91.523	-17.343	1.00	25.30	T4
	ATOM	4364	O	GLU	A	117	35.438	91.917	-16.661	1.00	34.00	T4
	ATOM	4365	N	LEU	A	118	33.347	91.104	-16.824	1.00	30.04	T4
	ATOM	4366	CA	LEU	A	118	33.138	91.039	-15.386	1.00	43.15	T4
	ATOM	4367	CB	LEU	A	118	31.834	91.727	-14.988	1.00	27.96	T4
30	ATOM	4368	CG	LEU	A	118	31.736	93.229	-15.212	1.00	35.83	T4
	ATOM	4369	CD1	LEU	A	118	30.411	93.716	-14.696	1.00	30.55	T4
	ATOM	4370	CD2	LEU	A	118	32.861	93.916	-14.496	1.00	37.58	T4
	ATOM	4371	C	LEU	A	118	33.061	89.579	-14.992	1.00	30.55	T4
	ATOM	4372	O	LEU	A	118	32.569	88.759	-15.756	1.00	27.33	T4
35	ATOM	4373	N	GLN	A	119	33.553	89.258	-13.803	1.00	37.93	T4
	ATOM	4374	CA	GLN	A	119	33.515	87.888	-13.316	1.00	29.50	T4
	ATOM	4375	CB	GLN	A	119	34.734	87.122	-13.815	1.00	28.78	T4
	ATOM	4376	CG	GLN	A	119	36.051	87.661	-13.309	1.00	32.17	T4
	ATOM	4377	CD	GLN	A	119	37.237	86.929	-13.903	1.00	26.87	T4
40	ATOM	4378	OE1	GLN	A	119	38.340	87.007	-13.384	1.00	28.78	T4
	ATOM	4379	NE2	GLN	A	119	37.014	86.226	-15.007	1.00	30.11	T4
	ATOM	4380	C	GLN	A	119	33.448	87.837	-11.791	1.00	33.53	T4
	ATOM	4381	O	GLN	A	119	33.871	88.769	-11.113	1.00	30.22	T4
	ATOM	4382	N	LEU	A	120	32.904	86.742	-11.268	1.00	31.61	T4
45	ATOM	4383	CA	LEU	A	120	32.744	86.529	-9.831	1.00	29.77	T4
	ATOM	4384	CB	LEU	A	120	31.303	86.097	-9.553	1.00	37.15	T4
	ATOM	4385	CG	LEU	A	120	30.661	86.029	-8.163	1.00	28.63	T4
	ATOM	4386	CD1	LEU	A	120	31.586	85.351	-7.179	1.00	35.10	T4
	ATOM	4387	CD2	LEU	A	120	30.315	87.422	-7.701	1.00	29.77	T4
50	ATOM	4388	C	LEU	A	120	33.702	85.409	-9.447	1.00	32.01	T4
	ATOM	4389	O	LEU	A	120	33.602	84.303	-9.970	1.00	35.25	T4
	ATOM	4390	N	ALA	A	121	34.624	85.678	-8.530	1.00	27.87	T4
	ATOM	4391	CA	ALA	A	121	35.589	84.661	-8.143	1.00	24.34	T4
	ATOM	4392	CB	ALA	A	121	36.953	85.017	-8.724	1.00	26.83	T4
55	ATOM	4393	C	ALA	A	121	35.710	84.413	-6.644	1.00	30.67	T4
	ATOM	4394	O	ALA	A	121	35.650	85.336	-5.835	1.00	27.10	T4
	ATOM	4395	N	ILE	A	122	35.876	83.144	-6.288	1.00	24.77	T4
	ATOM	4396	CA	ILE	A	122	36.057	82.740	-4.897	1.00	28.55	T4
	ATOM	4397	CB	ILE	A	122	35.151	81.565	-4.531	1.00	28.75	T4
60	ATOM	4398	CG2	ILE	A	122	35.331	81.227	-3.066	1.00	32.79	T4
	ATOM	4399	CG1	ILE	A	122	33.693	81.934	-4.808	1.00	31.01	T4
	ATOM	4400	CD1	ILE	A	122	32.709	80.854	-4.466	1.00	39.97	T4
	ATOM	4401	C	ILE	A	122	37.514	82.324	-4.695	1.00	30.41	T4
	ATOM	4402	O	ILE	A	122	37.994	81.374	-5.308	1.00	32.76	T4
65	ATOM	4403	N	PRO	A	123	38.238	83.042	-3.830	1.00	32.85	T4
	ATOM	4404	CD	PRO	A	123	37.778	84.254	-3.139	1.00	31.08	T4

	ATOM	4405	CA	PRO	A	123	39.649	82.783	-3.524	1.00	33.02	T4
	ATOM	4406	CB	PRO	A	123	40.078	84.038	-2.754	1.00	31.17	T4
	ATOM	4407	CG	PRO	A	123	39.035	85.061	-3.106	1.00	25.47	T4
	ATOM	4408	C	PRO	A	123	39.885	81.516	-2.696	1.00	33.00	T4
5	ATOM	4409	O	PRO	A	123	40.490	81.573	-1.624	1.00	31.83	T4
	ATOM	4410	N	ARG	A	124	39.392	80.383	-3.180	1.00	27.08	T4
	ATOM	4411	CA	ARG	A	124	39.576	79.110	-2.494	1.00	28.43	T4
	ATOM	4412	CB	ARG	A	124	38.435	78.818	-1.545	1.00	34.92	T4
	ATOM	4413	CG	ARG	A	124	38.638	79.426	-0.200	1.00	35.98	T4
10	ATOM	4414	CD	ARG	A	124	38.254	78.422	0.866	1.00	34.80	T4
	ATOM	4415	NE	ARG	A	124	39.238	77.356	0.999	1.00	25.96	T4
	ATOM	4416	CZ	ARG	A	124	39.047	76.272	1.742	1.00	31.21	T4
	ATOM	4417	NH1	ARG	A	124	37.910	76.123	2.399	1.00	25.02	T4
	ATOM	4418	NH2	ARG	A	124	39.990	75.344	1.840	1.00	27.95	T4
15	ATOM	4419	C	ARG	A	124	39.654	78.012	-3.517	1.00	40.33	T4
	ATOM	4420	O	ARG	A	124	39.143	78.165	-4.629	1.00	36.19	T4
	ATOM	4421	N	GLU	A	125	40.278	76.899	-3.144	1.00	25.95	T4
	ATOM	4422	CA	GLU	A	125	40.429	75.804	-4.087	1.00	30.60	T4
	ATOM	4423	CB	GLU	A	125	41.525	74.852	-3.629	1.00	29.06	T4
20	ATOM	4424	CG	GLU	A	125	42.911	75.458	-3.795	1.00	31.66	T4
	ATOM	4425	CD	GLU	A	125	43.939	74.450	-4.316	1.00	27.50	T4
	ATOM	4426	OE1	GLU	A	125	44.102	73.377	-3.680	1.00	26.68	T4
	ATOM	4427	OE2	GLU	A	125	44.589	74.734	-5.359	1.00	30.62	T4
	ATOM	4428	C	GLU	A	125	39.145	75.040	-4.394	1.00	32.67	T4
25	ATOM	4429	O	GLU	A	125	38.815	74.830	-5.563	1.00	26.24	T4
	ATOM	4430	N	ASN	A	126	38.417	74.604	-3.378	1.00	31.46	T4
	ATOM	4431	CA	ASN	A	126	37.173	73.909	-3.671	1.00	27.17	T4
	ATOM	4432	CB	ASN	A	126	37.331	72.397	-3.546	1.00	27.06	T4
	ATOM	4433	CG	ASN	A	126	37.942	71.776	-4.795	1.00	33.52	T4
30	ATOM	4434	OD1	ASN	A	126	39.159	71.835	-5.013	1.00	33.20	T4
	ATOM	4435	ND2	ASN	A	126	37.093	71.192	-5.637	1.00	34.74	T4
	ATOM	4436	C	ASN	A	126	36.123	74.406	-2.724	1.00	29.57	T4
	ATOM	4437	O	ASN	A	126	35.594	73.661	-1.894	1.00	31.71	T4
	ATOM	4438	N	ALA	A	127	35.832	75.693	-2.856	1.00	28.70	T4
35	ATOM	4439	CA	ALA	A	127	34.863	76.347	-2.004	1.00	31.18	T4
	ATOM	4440	CB	ALA	A	127	34.568	77.728	-2.556	1.00	27.65	T4
	ATOM	4441	C	ALA	A	127	33.576	75.547	-1.879	1.00	33.91	T4
	ATOM	4442	O	ALA	A	127	33.033	75.072	-2.877	1.00	28.93	T4
	ATOM	4443	N	GLN	A	128	33.107	75.381	-0.647	1.00	31.43	T4
40	ATOM	4444	CA	GLN	A	128	31.852	74.683	-0.404	1.00	37.59	T4
	ATOM	4445	CB	GLN	A	128	31.839	74.102	1.008	1.00	28.23	T4
	ATOM	4446	CG	GLN	A	128	32.859	72.984	1.184	1.00	25.23	T4
	ATOM	4447	CD	GLN	A	128	32.832	71.988	0.020	1.00	29.72	T4
	ATOM	4448	OE1	GLN	A	128	31.807	71.349	-0.257	1.00	32.75	T4
45	ATOM	4449	NE2	GLN	A	128	33.962	71.861	-0.670	1.00	30.03	T4
	ATOM	4450	C	GLN	A	128	30.755	75.731	-0.577	1.00	26.75	T4
	ATOM	4451	O	GLN	A	128	30.479	76.531	0.321	1.00	37.21	T4
	ATOM	4452	N	ILE	A	129	30.133	75.708	-1.746	1.00	36.17	T4
	ATOM	4453	CA	ILE	A	129	29.125	76.682	-2.125	1.00	36.56	T4
50	ATOM	4454	CB	ILE	A	129	29.686	77.471	-3.358	1.00	31.13	T4
	ATOM	4455	CG2	ILE	A	129	28.706	77.511	-4.514	1.00	35.86	T4
	ATOM	4456	CG1	ILE	A	129	30.135	78.848	-2.902	1.00	31.18	T4
	ATOM	4457	CD1	ILE	A	129	31.213	78.791	-1.847	1.00	28.78	T4
	ATOM	4458	C	ILE	A	129	27.763	76.076	-2.445	1.00	36.71	T4
55	ATOM	4459	O	ILE	A	129	27.662	74.887	-2.733	1.00	30.96	T4
	ATOM	4460	N	SER	A	130	26.714	76.887	-2.374	1.00	29.71	T4
	ATOM	4461	CA	SER	A	130	25.380	76.418	-2.729	1.00	28.89	T4
	ATOM	4462	CB	SER	A	130	24.333	76.967	-1.770	1.00	27.29	T4
	ATOM	4463	OG	SER	A	130	23.031	76.719	-2.273	1.00	32.84	T4
60	ATOM	4464	C	SER	A	130	25.092	76.944	-4.133	1.00	27.13	T4
	ATOM	4465	O	SER	A	130	25.216	78.144	-4.378	1.00	28.97	T4
	ATOM	4466	N	LEU	A	131	24.715	76.065	-5.057	1.00	27.41	T4
	ATOM	4467	CA	LEU	A	131	24.435	76.509	-6.421	1.00	30.22	T4
	ATOM	4468	CB	LEU	A	131	25.025	75.530	-7.435	1.00	28.78	T4
65	ATOM	4469	CG	LEU	A	131	26.523	75.661	-7.686	1.00	25.36	T4
	ATOM	4470	CD1	LEU	A	131	27.285	75.455	-6.414	1.00	27.76	T4

	ATOM	4471	CD2	LEU	A	131	26.938	74.639	-8.705	1.00	38.01	T4
	ATOM	4472	C	LEU	A	131	22.957	76.732	-6.731	1.00	31.88	T4
	ATOM	4473	O	LEU	A	131	22.515	76.531	-7.863	1.00	28.30	T4
	ATOM	4474	N	ASP	A	132	22.193	77.160	-5.731	1.00	25.81	T4
5	ATOM	4475	CA	ASP	A	132	20.776	77.420	-5.927	1.00	26.82	T4
	ATOM	4476	CB	ASP	A	132	20.018	77.237	-4.612	1.00	35.55	T4
	ATOM	4477	CG	ASP	A	132	19.686	75.776	-4.328	1.00	31.58	T4
	ATOM	4478	OD1	ASP	A	132	19.262	75.465	-3.185	1.00	28.62	T4
	ATOM	4479	OD2	ASP	A	132	19.839	74.943	-5.253	1.00	38.39	T4
10	ATOM	4480	C	ASP	A	132	20.519	78.817	-6.488	1.00	32.00	T4
	ATOM	4481	O	ASP	A	132	21.176	79.791	-6.112	1.00	29.44	T4
	ATOM	4482	N	GLY	A	133	19.556	78.898	-7.399	1.00	27.95	T4
	ATOM	4483	CA	GLY	A	133	19.203	80.162	-8.008	1.00	39.06	T4
	ATOM	4484	C	GLY	A	133	18.888	81.293	-7.048	1.00	28.00	T4
15	ATOM	4485	O	GLY	A	133	19.149	82.437	-7.387	1.00	30.59	T4
	ATOM	4486	N	ASP	A	134	18.336	80.995	-5.869	1.00	30.16	T4
	ATOM	4487	CA	ASP	A	134	18.006	82.040	-4.902	1.00	32.10	T4
	ATOM	4488	CB	ASP	A	134	17.210	81.532	-3.704	1.00	30.54	T4
	ATOM	4489	CG	ASP	A	134	16.434	80.316	-3.990	1.00	33.91	T4
20	ATOM	4490	OD1	ASP	A	134	15.533	80.402	-4.836	1.00	29.24	T4
	ATOM	4491	OD2	ASP	A	134	16.719	79.280	-3.361	1.00	14.25	T4
	ATOM	4492	C	ASP	A	134	19.242	82.614	-4.277	1.00	34.91	T4
	ATOM	4493	O	ASP	A	134	19.481	83.808	-4.313	1.00	23.95	T4
	ATOM	4494	N	VAL	A	135	20.007	81.734	-3.659	1.00	30.57	T4
25	ATOM	4495	CA	VAL	A	135	21.182	82.138	-2.933	1.00	26.30	T4
	ATOM	4496	CB	VAL	A	135	21.605	80.998	-2.040	1.00	29.58	T4
	ATOM	4497	CG1	VAL	A	135	20.477	80.692	-1.066	1.00	27.69	T4
	ATOM	4498	CG2	VAL	A	135	21.921	79.783	-2.890	1.00	32.79	T4
	ATOM	4499	C	VAL	A	135	22.399	82.707	-3.653	1.00	27.29	T4
30	ATOM	4500	O	VAL	A	135	23.039	83.611	-3.122	1.00	30.02	T4
	ATOM	4501	N	THR	A	136	22.750	82.208	-4.833	1.00	35.79	T4
	ATOM	4502	CA	THR	A	136	23.920	82.780	-5.500	1.00	31.82	T4
	ATOM	4503	CB	THR	A	136	25.133	81.796	-5.468	1.00	27.80	T4
	ATOM	4504	OG1	THR	A	136	24.971	80.787	-6.453	1.00	29.16	T4
35	ATOM	4505	CG2	THR	A	136	25.225	81.115	-4.122	1.00	27.68	T4
	ATOM	4506	C	THR	A	136	23.643	83.262	-6.933	1.00	24.23	T4
	ATOM	4507	O	THR	A	136	23.231	82.496	-7.803	1.00	27.75	T4
	ATOM	4508	N	PHE	A	137	23.873	84.553	-7.160	1.00	36.22	T4
	ATOM	4509	CA	PHE	A	137	23.631	85.165	-8.458	1.00	29.04	T4
40	ATOM	4510	CB	PHE	A	137	22.203	85.700	-8.489	1.00	32.24	T4
	ATOM	4511	CG	PHE	A	137	21.795	86.423	-7.236	1.00	20.81	T4
	ATOM	4512	CD1	PHE	A	137	22.153	87.743	-7.030	1.00	35.68	T4
	ATOM	4513	CD2	PHE	A	137	21.043	85.785	-6.270	1.00	31.72	T4
	ATOM	4514	CE1	PHE	A	137	21.765	88.413	-5.880	1.00	27.81	T4
45	ATOM	4515	CE2	PHE	A	137	20.652	86.447	-5.119	1.00	36.83	T4
	ATOM	4516	CZ	PHE	A	137	21.012	87.760	-4.925	1.00	26.95	T4
	ATOM	4517	C	PHE	A	137	24.632	86.275	-8.780	1.00	30.24	T4
	ATOM	4518	O	PHE	A	137	25.325	86.763	-7.893	1.00	31.64	T4
	ATOM	4519	N	PHE	A	138	24.695	86.680	-10.047	1.00	32.75	T4
50	ATOM	4520	CA	PHE	A	138	25.643	87.700	-10.473	1.00	35.61	T4
	ATOM	4521	CB	PHE	A	138	26.723	87.012	-11.306	1.00	30.91	T4
	ATOM	4522	CG	PHE	A	138	27.940	87.845	-11.552	1.00	32.88	T4
	ATOM	4523	CD1	PHE	A	138	28.269	88.897	-10.713	1.00	27.90	T4
	ATOM	4524	CD2	PHE	A	138	28.760	87.577	-12.640	1.00	28.57	T4
55	ATOM	4525	CE1	PHE	A	138	29.395	89.676	-10.960	1.00	36.79	T4
	ATOM	4526	CE2	PHE	A	138	29.886	88.346	-12.895	1.00	33.89	T4
	ATOM	4527	CZ	PHE	A	138	30.205	89.398	-12.057	1.00	27.25	T4
	ATOM	4528	C	PHE	A	138	24.967	88.861	-11.233	1.00	29.90	T4
	ATOM	4529	O	PHE	A	138	24.196	88.647	-12.173	1.00	30.04	T4
60	ATOM	4530	N	GLY	A	139	25.305	90.086	-10.809	1.00	25.74	T4
	ATOM	4531	CA	GLY	A	139	24.722	91.329	-11.313	1.00	26.69	T4
	ATOM	4532	C	GLY	A	139	24.944	91.982	-12.655	1.00	26.81	T4
	ATOM	4533	O	GLY	A	139	24.869	91.316	-13.666	1.00	34.52	T4
	ATOM	4534	N	ALA	A	140	25.147	93.304	-12.641	1.00	28.73	T4
65	ATOM	4535	CA	ALA	A	140	25.376	94.151	-13.830	1.00	26.92	T4
	ATOM	4536	CB	ALA	A	140	26.169	93.401	-14.869	1.00	32.99	T4

	ATOM	4537	C	ALA	A	140	24.146	94.779	-14.493	1.00	27.60	T4
	ATOM	4538	O	ALA	A	140	23.389	94.113	-15.185	1.00	28.35	T4
	ATOM	4539	N	LEU	A	141	23.982	96.083	-14.285	1.00	33.47	T4
	ATOM	4540	CA	LEU	A	141	22.875	96.871	-14.838	1.00	29.14	T4
5	ATOM	4541	CB	LEU	A	141	21.786	97.074	-13.775	1.00	33.42	T4
	ATOM	4542	CG	LEU	A	141	20.620	98.024	-14.076	1.00	32.11	T4
	ATOM	4543	CD1	LEU	A	141	19.475	97.759	-13.135	1.00	33.35	T4
	ATOM	4544	CD2	LEU	A	141	21.077	99.457	-13.937	1.00	31.38	T4
	ATOM	4545	C	LEU	A	141	23.432	98.226	-15.271	1.00	29.36	T4
10	ATOM	4546	O	LEU	A	141	24.171	98.856	-14.528	1.00	28.62	T4
	ATOM	4547	N	LYS	A	142	23.067	98.688	-16.456	1.00	35.38	T4
	ATOM	4548	CA	LYS	A	142	23.589	99.957	-16.940	1.00	31.90	T4
	ATOM	4549	CB	LYS	A	142	23.705	99.925	-18.463	1.00	36.04	T4
	ATOM	4550	CG	LYS	A	142	24.241	101.219	-19.029	1.00	24.73	T4
15	ATOM	4551	CD	LYS	A	142	24.631	101.110	-20.484	1.00	24.59	T4
	ATOM	4552	CE	LYS	A	142	25.192	102.441	-20.975	1.00	38.39	T4
	ATOM	4553	NZ	LYS	A	142	25.551	102.390	-22.422	1.00	36.04	T4
	ATOM	4554	C	LYS	A	142	22.808	101.197	-16.520	1.00	35.77	T4
	ATOM	4555	O	LYS	A	142	21.596	101.264	-16.689	1.00	31.95	T4
20	ATOM	4556	N	LEU	A	143	23.518	102.185	-15.980	1.00	33.69	T4
	ATOM	4557	CA	LEU	A	143	22.900	103.439	-15.548	1.00	26.78	T4
	ATOM	4558	CB	LEU	A	143	23.777	104.128	-14.504	1.00	41.17	T4
	ATOM	4559	CG	LEU	A	143	24.131	103.365	-13.223	1.00	34.12	T4
	ATOM	4560	CD1	LEU	A	143	25.085	104.205	-12.388	1.00	38.65	T4
25	ATOM	4561	CD2	LEU	A	143	22.873	103.056	-12.429	1.00	33.84	T4
	ATOM	4562	C	LEU	A	143	22.707	104.385	-16.735	1.00	25.82	T4
	ATOM	4563	O	LEU	A	143	23.412	104.292	-17.737	1.00	33.19	T4
	ATOM	4564	N	LEU	A	144	21.749	105.297	-16.624	1.00	31.38	T4
	ATOM	4565	CA	LEU	A	144	21.501	106.248	-17.697	1.00	30.56	T4
30	ATOM	4566	CB	LEU	A	144	20.040	106.688	-17.696	1.00	28.74	T4
	ATOM	4567	CG	LEU	A	144	19.024	105.600	-18.032	1.00	34.96	T4
	ATOM	4568	CD1	LEU	A	144	17.631	106.147	-17.811	1.00	29.78	T4
	ATOM	4569	CD2	LEU	A	144	19.199	105.131	-19.473	1.00	30.17	T4
	ATOM	4570	C	LEU	A	144	22.384	107.468	-17.520	1.00	32.43	T4
35	ATOM	4571	O	LEU	A	144	22.892	107.668	-16.394	1.00	29.54	T4
	ATOM	4572	OXT	LEU	A	144	22.538	108.217	-18.507	1.00	31.71	T4
	ATOM	4573	CB	VAL	A	1	19.493	113.420	-19.948	1.00	34.35	T5
	ATOM	4574	CG1	VAL	A	1	20.191	112.650	-21.095	1.00	35.26	T5
	ATOM	4575	CG2	VAL	A	1	18.626	114.545	-20.517	1.00	28.30	T5
40	ATOM	4576	C	VAL	A	1	19.534	111.393	-18.421	1.00	31.46	T5
	ATOM	4577	O	VAL	A	1	19.396	110.195	-18.686	1.00	31.94	T5
	ATOM	4578	N	VAL	A	1	17.808	113.195	-18.082	1.00	27.44	T5
	ATOM	4579	CA	VAL	A	1	18.616	112.444	-19.086	1.00	30.58	T5
	ATOM	4580	N	THR	A	2	20.461	111.833	-17.567	1.00	28.08	T5
45	ATOM	4581	CA	THR	A	2	21.378	110.900	-16.905	1.00	32.44	T5
	ATOM	4582	CB	THR	A	2	22.851	111.290	-17.130	1.00	33.78	T5
	ATOM	4583	OG1	THR	A	2	23.150	112.467	-16.373	1.00	36.09	T5
	ATOM	4584	CG2	THR	A	2	23.118	111.548	-18.600	1.00	27.63	T5
	ATOM	4585	C	THR	A	2	21.158	110.797	-15.392	1.00	28.21	T5
50	ATOM	4586	O	THR	A	2	20.410	111.575	-14.808	1.00	23.48	T5
	ATOM	4587	N	GLN	A	3	21.832	109.835	-14.765	1.00	26.60	T5
	ATOM	4588	CA	GLN	A	3	21.710	109.626	-13.329	1.00	35.34	T5
	ATOM	4589	CB	GLN	A	3	21.372	108.166	-13.044	1.00	25.90	T5
	ATOM	4590	CG	GLN	A	3	20.275	107.608	-13.897	1.00	32.51	T5
55	ATOM	4591	CD	GLN	A	3	19.865	106.222	-13.453	1.00	24.71	T5
	ATOM	4592	OE1	GLN	A	3	19.411	106.039	-12.327	1.00	30.67	T5
	ATOM	4593	NE2	GLN	A	3	20.022	105.237	-14.337	1.00	34.62	T5
	ATOM	4594	C	GLN	A	3	23.002	109.979	-12.590	1.00	30.26	T5
	ATOM	4595	O	GLN	A	3	23.973	109.212	-12.623	1.00	35.21	T5
60	ATOM	4596	N	ASP	A	4	23.020	111.123	-11.915	1.00	32.99	T5
	ATOM	4597	CA	ASP	A	4	24.216	111.510	-11.188	1.00	31.78	T5
	ATOM	4598	CB	ASP	A	4	24.036	112.885	-10.541	1.00	36.72	T5
	ATOM	4599	CG	ASP	A	4	23.860	113.990	-11.561	1.00	28.71	T5
	ATOM	4600	OD1	ASP	A	4	24.197	113.755	-12.745	1.00	27.30	T5
65	ATOM	4601	OD2	ASP	A	4	23.400	115.093	-11.183	1.00	29.56	T5
	ATOM	4602	C	ASP	A	4	24.509	110.478	-10.113	1.00	33.47	T5

	ATOM	4603	O	ASP	A	4	23.587	109.864	-9.569	1.00	27.57	T5
	ATOM	4604	N	CYS	A	5	25.795	110.281	-9.825	1.00	29.52	T5
	ATOM	4605	CA	CYS	A	5	26.232	109.339	-8.795	1.00	23.67	T5
	ATOM	4606	CB	CYS	A	5	26.241	107.895	-9.333	1.00	27.60	T5
5	ATOM	4607	SG	CYS	A	5	26.573	107.697	-11.122	1.00	27.12	T5
	ATOM	4608	C	CYS	A	5	27.614	109.721	-8.296	1.00	26.79	T5
	ATOM	4609	O	CYS	A	5	28.412	110.254	-9.045	1.00	25.75	T5
	ATOM	4610	N	LEU	A	6	27.884	109.479	-7.019	1.00	33.09	T5
	ATOM	4611	CA	LEU	A	6	29.188	109.781	-6.435	1.00	28.46	T5
10	ATOM	4612	CB	LEU	A	6	29.127	111.071	-5.629	1.00	32.64	T5
	ATOM	4613	CG	LEU	A	6	30.404	111.492	-4.908	1.00	25.38	T5
	ATOM	4614	CD1	LEU	A	6	30.465	113.004	-4.796	1.00	30.82	T5
	ATOM	4615	CD2	LEU	A	6	30.447	110.860	-3.544	1.00	29.90	T5
	ATOM	4616	C	LEU	A	6	29.554	108.619	-5.535	1.00	35.48	T5
15	ATOM	4617	O	LEU	A	6	28.732	108.159	-4.757	1.00	27.61	T5
	ATOM	4618	N	GLN	A	7	30.783	108.134	-5.642	1.00	29.89	T5
	ATOM	4619	CA	GLN	A	7	31.206	107.004	-4.833	1.00	25.96	T5
	ATOM	4620	CB	GLN	A	7	31.351	105.775	-5.713	1.00	37.07	T5
	ATOM	4621	CG	GLN	A	7	31.524	104.480	-4.958	1.00	30.75	T5
20	ATOM	4622	CD	GLN	A	7	31.422	103.288	-5.882	1.00	35.17	T5
	ATOM	4623	OE1	GLN	A	7	32.287	103.060	-6.714	1.00	29.56	T5
	ATOM	4624	NE2	GLN	A	7	30.351	102.535	-5.754	1.00	28.68	T5
	ATOM	4625	C	GLN	A	7	32.510	107.268	-4.115	1.00	21.35	T5
	ATOM	4626	O	GLN	A	7	33.417	107.884	-4.667	1.00	32.88	T5
25	ATOM	4627	N	LEU	A	8	32.591	106.796	-2.876	1.00	33.27	T5
	ATOM	4628	CA	LEU	A	8	33.781	106.975	-2.053	1.00	28.85	T5
	ATOM	4629	CB	LEU	A	8	33.423	107.707	-0.757	1.00	34.74	T5
	ATOM	4630	CG	LEU	A	8	33.392	109.242	-0.723	1.00	28.20	T5
	ATOM	4631	CD1	LEU	A	8	33.392	109.834	-2.118	1.00	28.57	T5
30	ATOM	4632	CD2	LEU	A	8	32.173	109.680	0.054	1.00	34.63	T5
	ATOM	4633	C	LEU	A	8	34.447	105.647	-1.719	1.00	30.37	T5
	ATOM	4634	O	LEU	A	8	33.796	104.608	-1.664	1.00	36.40	T5
	ATOM	4635	N	ILE	A	9	35.754	105.703	-1.491	1.00	34.73	T5
	ATOM	4636	CA	ILE	A	9	36.563	104.529	-1.170	1.00	22.90	T5
35	ATOM	4637	CB	ILE	A	9	37.550	104.238	-2.303	1.00	33.78	T5
	ATOM	4638	CG2	ILE	A	9	38.666	103.371	-1.813	1.00	32.92	T5
	ATOM	4639	CG1	ILE	A	9	36.850	103.560	-3.458	1.00	31.58	T5
	ATOM	4640	CD1	ILE	A	9	37.801	103.300	-4.587	1.00	38.61	T5
	ATOM	4641	C	ILE	A	9	37.385	104.806	0.084	1.00	31.12	T5
40	ATOM	4642	O	ILE	A	9	37.843	105.926	0.290	1.00	30.06	T5
	ATOM	4643	N	ALA	A	10	37.591	103.793	0.914	1.00	29.93	T5
	ATOM	4644	CA	ALA	A	10	38.381	103.994	2.121	1.00	31.67	T5
	ATOM	4645	CB	ALA	A	10	38.406	102.730	2.950	1.00	27.10	T5
	ATOM	4646	C	ALA	A	10	39.809	104.407	1.770	1.00	33.78	T5
45	ATOM	4647	O	ALA	A	10	40.457	103.797	0.916	1.00	26.62	T5
	ATOM	4648	N	ASP	A	11	40.294	105.451	2.433	1.00	30.64	T5
	ATOM	4649	CA	ASP	A	11	41.648	105.934	2.205	1.00	29.66	T5
	ATOM	4650	CB	ASP	A	11	41.718	107.448	2.366	1.00	34.30	T5
	ATOM	4651	CG	ASP	A	11	43.128	107.967	2.261	1.00	36.53	T5
50	ATOM	4652	OD1	ASP	A	11	43.881	107.428	1.427	1.00	29.89	T5
	ATOM	4653	OD2	ASP	A	11	43.485	108.910	2.998	1.00	33.05	T5
	ATOM	4654	C	ASP	A	11	42.596	105.272	3.183	1.00	27.04	T5
	ATOM	4655	O	ASP	A	11	42.827	105.764	4.282	1.00	33.43	T5
	ATOM	4656	N	SER	A	12	43.140	104.144	2.749	1.00	33.38	T5
55	ATOM	4657	CA	SER	A	12	44.054	103.336	3.540	1.00	25.46	T5
	ATOM	4658	CB	SER	A	12	44.377	102.033	2.794	1.00	28.98	T5
	ATOM	4659	OG	SER	A	12	44.944	102.277	1.505	1.00	29.33	T5
	ATOM	4660	C	SER	A	12	45.346	104.042	3.876	1.00	26.08	T5
	ATOM	4661	O	SER	A	12	46.349	103.391	4.157	1.00	34.38	T5
60	ATOM	4662	N	GLU	A	13	45.345	105.369	3.845	1.00	32.06	T5
	ATOM	4663	CA	GLU	A	13	46.569	106.092	4.159	1.00	26.32	T5
	ATOM	4664	CB	GLU	A	13	47.274	106.507	2.886	1.00	29.91	T5
	ATOM	4665	CG	GLU	A	13	48.235	105.446	2.422	1.00	29.99	T5
	ATOM	4666	CD	GLU	A	13	49.005	105.890	1.204	1.00	31.18	T5
65	ATOM	4667	OE1	GLU	A	13	49.416	107.078	1.172	1.00	26.03	T5
	ATOM	4668	OE2	GLU	A	13	49.205	105.059	0.278	1.00	37.24	T5

	ATOM	4669	C	GLU	A	13	46.436	107.275	5.079	1.00	31.55	T5
	ATOM	4670	O	GLU	A	13	47.250	108.194	5.040	1.00	32.64	T5
	ATOM	4671	N	THR	A	14	45.396	107.250	5.900	1.00	28.65	T5
	ATOM	4672	CA	THR	A	14	45.168	108.295	6.883	1.00	34.82	T5
5	ATOM	4673	CB	THR	A	14	44.291	109.459	6.360	1.00	27.83	T5
	ATOM	4674	OG1	THR	A	14	43.036	108.947	5.924	1.00	32.56	T5
	ATOM	4675	CG2	THR	A	14	44.982	110.190	5.209	1.00	33.88	T5
	ATOM	4676	C	THR	A	14	44.461	107.606	8.028	1.00	29.73	T5
	ATOM	4677	O	THR	A	14	43.756	106.613	7.835	1.00	27.29	T5
10	ATOM	4678	N	PRO	A	15	44.657	108.113	9.245	1.00	30.89	T5
	ATOM	4679	CD	PRO	A	15	45.455	109.301	9.579	1.00	31.07	T5
	ATOM	4680	CA	PRO	A	15	44.046	107.545	10.444	1.00	36.92	T5
	ATOM	4681	CB	PRO	A	15	44.598	108.428	11.565	1.00	31.16	T5
	ATOM	4682	CG	PRO	A	15	45.861	109.003	10.981	1.00	31.19	T5
15	ATOM	4683	C	PRO	A	15	42.537	107.611	10.380	1.00	31.20	T5
	ATOM	4684	O	PRO	A	15	41.985	108.549	9.838	1.00	29.58	T5
	ATOM	4685	N	THR	A	16	41.873	106.616	10.943	1.00	32.25	T5
	ATOM	4686	CA	THR	A	16	40.426	106.611	10.959	1.00	32.62	T5
	ATOM	4687	CB	THR	A	16	39.896	105.253	11.411	1.00	26.35	T5
20	ATOM	4688	OG1	THR	A	16	40.176	105.061	12.806	1.00	29.80	T5
	ATOM	4689	CG2	THR	A	16	40.573	104.156	10.625	1.00	27.62	T5
	ATOM	4690	C	THR	A	16	39.967	107.680	11.949	1.00	30.28	T5
	ATOM	4691	O	THR	A	16	40.236	107.582	13.146	1.00	34.94	T5
	ATOM	4692	N	ILE	A	17	39.270	108.696	11.448	1.00	30.33	T5
25	ATOM	4693	CA	ILE	A	17	38.775	109.787	12.289	1.00	31.99	T5
	ATOM	4694	CB	ILE	A	17	37.886	110.726	11.480	1.00	32.32	T5
	ATOM	4695	CG2	ILE	A	17	37.375	111.836	12.361	1.00	30.96	T5
	ATOM	4696	CG1	ILE	A	17	38.688	111.302	10.315	1.00	31.80	T5
	ATOM	4697	CD1	ILE	A	17	37.900	112.199	9.403	1.00	25.80	T5
30	ATOM	4698	C	ILE	A	17	37.998	109.358	13.541	1.00	25.78	T5
	ATOM	4699	O	ILE	A	17	37.087	108.528	13.477	1.00	29.93	T5
	ATOM	4700	N	GLN	A	18	38.370	109.937	14.679	1.00	29.60	T5
	ATOM	4701	CA	GLN	A	18	37.717	109.640	15.949	1.00	32.88	T5
	ATOM	4702	CB	GLN	A	18	38.731	109.188	16.994	1.00	30.24	T5
35	ATOM	4703	CG	GLN	A	18	38.383	107.856	17.586	1.00	34.53	T5
	ATOM	4704	CD	GLN	A	18	38.641	106.740	16.608	1.00	33.74	T5
	ATOM	4705	OE1	GLN	A	18	39.790	106.345	16.389	1.00	23.29	T5
	ATOM	4706	NE2	GLN	A	18	37.581	106.230	15.996	1.00	27.09	T5
	ATOM	4707	C	GLN	A	18	36.992	110.872	16.466	1.00	28.94	T5
40	ATOM	4708	O	GLN	A	18	37.537	111.976	16.463	1.00	29.74	T5
	ATOM	4709	N	LYS	A	19	35.758	110.680	16.919	1.00	19.43	T5
	ATOM	4710	CA	LYS	A	19	34.964	111.794	17.430	1.00	34.07	T5
	ATOM	4711	CB	LYS	A	19	34.661	112.761	16.295	1.00	32.06	T5
	ATOM	4712	CG	LYS	A	19	33.663	113.824	16.657	1.00	31.51	T5
45	ATOM	4713	CD	LYS	A	19	33.576	114.874	15.557	1.00	31.62	T5
	ATOM	4714	CE	LYS	A	19	32.540	115.953	15.886	1.00	30.27	T5
	ATOM	4715	NZ	LYS	A	19	32.469	117.030	14.842	1.00	35.19	T5
	ATOM	4716	C	LYS	A	19	33.659	111.354	18.089	1.00	29.73	T5
	ATOM	4717	O	LYS	A	19	32.897	110.583	17.513	1.00	28.58	T5
50	ATOM	4718	N	GLY	A	20	33.410	111.863	19.295	1.00	31.62	T5
	ATOM	4719	CA	GLY	A	20	32.208	111.505	20.026	1.00	30.03	T5
	ATOM	4720	C	GLY	A	20	32.098	110.003	20.171	1.00	29.65	T5
	ATOM	4721	O	GLY	A	20	31.013	109.435	20.034	1.00	31.10	T5
	ATOM	4722	N	SER	A	21	33.229	109.363	20.452	1.00	28.31	T5
55	ATOM	4723	CA	SER	A	21	33.296	107.908	20.603	1.00	29.30	T5
	ATOM	4724	CB	SER	A	21	32.648	107.468	21.925	1.00	30.07	T5
	ATOM	4725	OG	SER	A	21	31.272	107.802	21.959	1.00	35.40	T5
	ATOM	4726	C	SER	A	21	32.657	107.160	19.416	1.00	26.12	T5
	ATOM	4727	O	SER	A	21	31.977	106.139	19.582	1.00	31.66	T5
60	ATOM	4728	N	TYR	A	22	32.902	107.694	18.223	1.00	24.60	T5
	ATOM	4729	CA	TYR	A	22	32.422	107.133	16.966	1.00	31.74	T5
	ATOM	4730	CB	TYR	A	22	31.393	108.061	16.331	1.00	36.33	T5
	ATOM	4731	CG	TYR	A	22	29.964	107.716	16.643	1.00	28.09	T5
	ATOM	4732	CD1	TYR	A	22	29.631	107.005	17.795	1.00	25.31	T5
65	ATOM	4733	CE1	TYR	A	22	28.296	106.702	18.097	1.00	25.54	T5
	ATOM	4734	CD2	TYR	A	22	28.931	108.122	15.799	1.00	36.67	T5

	ATOM	4735	CE2	TYR	A	22	27.597	107.829	16.093	1.00	29.97	T5
	ATOM	4736	CZ	TYR	A	22	27.290	107.116	17.243	1.00	30.81	T5
	ATOM	4737	OH	TYR	A	22	25.984	106.803	17.541	1.00	29.80	T5
	ATOM	4738	C	TYR	A	22	33.625	107.050	16.046	1.00	32.84	T5
5	ATOM	4739	O	TYR	A	22	34.519	107.896	16.116	1.00	33.67	T5
	ATOM	4740	N	THR	A	23	33.661	106.038	15.189	1.00	30.87	T5
	ATOM	4741	CA	THR	A	23	34.773	105.921	14.267	1.00	32.81	T5
	ATOM	4742	CB	THR	A	23	35.352	104.501	14.239	1.00	28.93	T5
	ATOM	4743	OG1	THR	A	23	35.401	103.967	15.569	1.00	34.05	T5
10	ATOM	4744	CG2	THR	A	23	36.767	104.543	13.684	1.00	30.10	T5
	ATOM	4745	C	THR	A	23	34.280	106.280	12.876	1.00	28.60	T5
	ATOM	4746	O	THR	A	23	33.247	105.780	12.432	1.00	32.14	T5
	ATOM	4747	N	PHE	A	24	35.010	107.160	12.198	1.00	36.61	T5
	ATOM	4748	CA	PHE	A	24	34.642	107.578	10.852	1.00	33.06	T5
15	ATOM	4749	CB	PHE	A	24	34.402	109.085	10.811	1.00	30.72	T5
	ATOM	4750	CG	PHE	A	24	33.218	109.526	11.613	1.00	29.51	T5
	ATOM	4751	CD1	PHE	A	24	33.311	109.679	12.992	1.00	38.02	T5
	ATOM	4752	CD2	PHE	A	24	31.994	109.760	10.997	1.00	26.89	T5
	ATOM	4753	CE1	PHE	A	24	32.197	110.058	13.744	1.00	31.11	T5
20	ATOM	4754	CE2	PHE	A	24	30.884	110.135	11.737	1.00	33.49	T5
	ATOM	4755	CZ	PHE	A	24	30.985	110.285	13.113	1.00	37.20	T5
	ATOM	4756	C	PHE	A	24	35.703	107.205	9.828	1.00	34.03	T5
	ATOM	4757	O	PHE	A	24	36.867	107.581	9.956	1.00	31.79	T5
	ATOM	4758	N	VAL	A	25	35.287	106.461	8.808	1.00	32.98	T5
25	ATOM	4759	CA	VAL	A	25	36.187	106.030	7.751	1.00	31.31	T5
	ATOM	4760	CB	VAL	A	25	35.478	105.086	6.780	1.00	32.77	T5
	ATOM	4761	CG1	VAL	A	25	36.411	104.706	5.643	1.00	28.31	T5
	ATOM	4762	CG2	VAL	A	25	35.000	103.865	7.516	1.00	27.94	T5
	ATOM	4763	C	VAL	A	25	36.708	107.217	6.956	1.00	26.52	T5
30	ATOM	4764	O	VAL	A	25	35.948	108.104	6.577	1.00	28.58	T5
	ATOM	4765	N	PRO	A	26	38.024	107.257	6.707	1.00	26.73	T5
	ATOM	4766	CD	PRO	A	26	39.044	106.371	7.287	1.00	28.93	T5
	ATOM	4767	CA	PRO	A	26	38.649	108.343	5.942	1.00	31.92	T5
	ATOM	4768	CB	PRO	A	26	40.137	108.157	6.225	1.00	30.44	T5
35	ATOM	4769	CG	PRO	A	26	40.177	107.316	7.478	1.00	28.01	T5
	ATOM	4770	C	PRO	A	26	38.334	108.096	4.470	1.00	30.18	T5
	ATOM	4771	O	PRO	A	26	38.840	107.143	3.886	1.00	34.61	T5
	ATOM	4772	N	TRP	A	27	37.511	108.937	3.861	1.00	28.52	T5
	ATOM	4773	CA	TRP	A	27	37.162	108.714	2.463	1.00	32.84	T5
40	ATOM	4774	CB	TRP	A	27	35.747	109.230	2.181	1.00	32.09	T5
	ATOM	4775	CG	TRP	A	27	34.699	108.515	2.946	1.00	31.47	T5
	ATOM	4776	CD2	TRP	A	27	34.480	107.102	2.988	1.00	28.13	T5
	ATOM	4777	CE2	TRP	A	27	33.428	106.871	3.894	1.00	26.62	T5
	ATOM	4778	CE3	TRP	A	27	35.075	106.006	2.353	1.00	30.31	T5
45	ATOM	4779	CD1	TRP	A	27	33.797	109.068	3.792	1.00	32.33	T5
	ATOM	4780	NE1	TRP	A	27	33.030	108.091	4.370	1.00	34.82	T5
	ATOM	4781	CZ2	TRP	A	27	32.953	105.586	4.183	1.00	26.73	T5
	ATOM	4782	CZ3	TRP	A	27	34.601	104.725	2.644	1.00	24.78	T5
	ATOM	4783	CH2	TRP	A	27	33.552	104.530	3.550	1.00	33.54	T5
50	ATOM	4784	C	TRP	A	27	38.121	109.278	1.419	1.00	31.14	T5
	ATOM	4785	O	TRP	A	27	38.971	110.127	1.698	1.00	30.89	T5
	ATOM	4786	N	LEU	A	28	37.964	108.772	0.203	1.00	30.49	T5
	ATOM	4787	CA	LEU	A	28	38.759	109.178	-0.935	1.00	33.82	T5
	ATOM	4788	CB	LEU	A	28	39.978	108.284	-1.053	1.00	37.07	T5
55	ATOM	4789	CG	LEU	A	28	41.115	108.863	-1.875	1.00	31.32	T5
	ATOM	4790	CD1	LEU	A	28	41.625	110.127	-1.189	1.00	22.42	T5
	ATOM	4791	CD2	LEU	A	28	42.218	107.822	-2.011	1.00	30.80	T5
	ATOM	4792	C	LEU	A	28	37.855	108.996	-2.150	1.00	31.81	T5
	ATOM	4793	O	LEU	A	28	37.259	107.940	-2.338	1.00	28.36	T5
60	ATOM	4794	N	LEU	A	29	37.742	110.029	-2.971	1.00	27.21	T5
	ATOM	4795	CA	LEU	A	29	36.876	109.951	-4.133	1.00	26.27	T5
	ATOM	4796	CB	LEU	A	29	36.971	111.225	-4.967	1.00	34.67	T5
	ATOM	4797	CG	LEU	A	29	36.092	111.185	-6.213	1.00	29.51	T5
	ATOM	4798	CD1	LEU	A	29	34.646	111.384	-5.821	1.00	27.68	T5
65	ATOM	4799	CD2	LEU	A	29	36.527	112.248	-7.180	1.00	25.88	T5
	ATOM	4800	C	LEU	A	29	37.191	108.774	-5.021	1.00	39.17	T5

	ATOM	4801	O	LEU	A	29	38.332	108.589	-5.429	1.00	26.04	T5
	ATOM	4802	N	SER	A	30	36.173	107.971	-5.302	1.00	26.92	T5
	ATOM	4803	CA	SER	A	30	36.322	106.840	-6.197	1.00	32.27	T5
	ATOM	4804	CB	SER	A	30	35.312	105.751	-5.869	1.00	36.35	T5
5	ATOM	4805	OG	SER	A	30	35.388	104.711	-6.819	1.00	30.61	T5
	ATOM	4806	C	SER	A	30	36.021	107.423	-7.572	1.00	27.48	T5
	ATOM	4807	O	SER	A	30	36.821	107.320	-8.496	1.00	29.27	T5
	ATOM	4808	N	PHE	A	31	34.860	108.054	-7.692	1.00	34.56	T5
	ATOM	4809	CA	PHE	A	31	34.457	108.682	-8.941	1.00	32.40	T5
10	ATOM	4810	CB	PHE	A	31	34.129	107.620	-9.997	1.00	31.66	T5
	ATOM	4811	CG	PHE	A	31	32.712	107.125	-9.948	1.00	33.98	T5
	ATOM	4812	CD1	PHE	A	31	31.699	107.815	-10.595	1.00	25.10	T5
	ATOM	4813	CD2	PHE	A	31	32.391	105.968	-9.259	1.00	30.26	T5
	ATOM	4814	CE1	PHE	A	31	30.388	107.359	-10.557	1.00	26.71	T5
15	ATOM	4815	CE2	PHE	A	31	31.082	105.506	-9.218	1.00	33.18	T5
	ATOM	4816	CZ	PHE	A	31	30.081	106.204	-9.869	1.00	24.57	T5
	ATOM	4817	C	PHE	A	31	33.242	109.556	-8.674	1.00	31.75	T5
	ATOM	4818	O	PHE	A	31	32.483	109.308	-7.745	1.00	29.75	T5
	ATOM	4819	N	LYS	A	32	33.071	110.587	-9.488	1.00	25.55	T5
20	ATOM	4820	CA	LYS	A	32	31.944	111.493	-9.348	1.00	29.30	T5
	ATOM	4821	CB	LYS	A	32	32.399	112.806	-8.738	1.00	33.49	T5
	ATOM	4822	CG	LYS	A	32	31.367	113.877	-8.851	1.00	33.25	T5
	ATOM	4823	CD	LYS	A	32	31.896	115.221	-8.425	1.00	37.05	T5
	ATOM	4824	CE	LYS	A	32	30.893	116.292	-8.815	1.00	25.74	T5
25	ATOM	4825	NZ	LYS	A	32	31.267	117.626	-8.275	1.00	34.28	T5
	ATOM	4826	C	LYS	A	32	31.356	111.746	-10.723	1.00	29.64	T5
	ATOM	4827	O	LYS	A	32	32.064	112.137	-11.647	1.00	29.05	T5
	ATOM	4828	N	ARG	A	33	30.057	111.524	-10.859	1.00	32.75	T5
	ATOM	4829	CA	ARG	A	33	29.382	111.701	-12.138	1.00	37.14	T5
30	ATOM	4830	CB	ARG	A	33	28.968	110.328	-12.676	1.00	31.89	T5
	ATOM	4831	CG	ARG	A	33	28.106	110.316	-13.919	1.00	29.81	T5
	ATOM	4832	CD	ARG	A	33	28.214	108.946	-14.569	1.00	37.49	T5
	ATOM	4833	NE	ARG	A	33	27.358	108.790	-15.738	1.00	29.08	T5
	ATOM	4834	CZ	ARG	A	33	26.070	108.482	-15.670	1.00	29.60	T5
35	ATOM	4835	NH1	ARG	A	33	25.501	108.292	-14.482	1.00	36.45	T5
	ATOM	4836	NH2	ARG	A	33	25.355	108.378	-16.786	1.00	29.91	T5
	ATOM	4837	C	ARG	A	33	28.171	112.601	-11.974	1.00	29.09	T5
	ATOM	4838	O	ARG	A	33	27.284	112.320	-11.168	1.00	33.58	T5
	ATOM	4839	N	GLY	A	34	28.144	113.691	-12.730	1.00	30.85	T5
40	ATOM	4840	CA	GLY	A	34	27.023	114.602	-12.647	1.00	28.97	T5
	ATOM	4841	C	GLY	A	34	27.239	115.737	-11.676	1.00	31.25	T5
	ATOM	4842	O	GLY	A	34	28.360	115.984	-11.234	1.00	33.44	T5
	ATOM	4843	N	SER	A	35	26.150	116.411	-11.320	1.00	33.96	T5
	ATOM	4844	CA	SER	A	35	26.222	117.554	-10.422	1.00	30.79	T5
45	ATOM	4845	CB	SER	A	35	25.723	118.788	-11.152	1.00	26.83	T5
	ATOM	4846	OG	SER	A	35	24.413	118.555	-11.651	1.00	32.46	T5
	ATOM	4847	C	SER	A	35	25.451	117.438	-9.119	1.00	26.16	T5
	ATOM	4848	O	SER	A	35	25.745	118.164	-8.175	1.00	28.82	T5
	ATOM	4849	N	ALA	A	36	24.471	116.539	-9.059	1.00	34.08	T5
50	ATOM	4850	CA	ALA	A	36	23.648	116.392	-7.862	1.00	30.69	T5
	ATOM	4851	CB	ALA	A	36	22.515	115.420	-8.132	1.00	27.39	T5
	ATOM	4852	C	ALA	A	36	24.359	116.002	-6.577	1.00	29.90	T5
	ATOM	4853	O	ALA	A	36	23.796	116.180	-5.502	1.00	33.38	T5
	ATOM	4854	N	LEU	A	37	25.583	115.483	-6.671	1.00	28.22	T5
55	ATOM	4855	CA	LEU	A	37	26.319	115.072	-5.474	1.00	27.94	T5
	ATOM	4856	CB	LEU	A	37	26.248	113.553	-5.329	1.00	33.72	T5
	ATOM	4857	CG	LEU	A	37	24.832	112.994	-5.189	1.00	26.59	T5
	ATOM	4858	CD1	LEU	A	37	24.817	111.518	-5.516	1.00	21.68	T5
	ATOM	4859	CD2	LEU	A	37	24.326	113.253	-3.790	1.00	27.99	T5
60	ATOM	4860	C	LEU	A	37	27.775	115.530	-5.468	1.00	30.92	T5
	ATOM	4861	O	LEU	A	37	28.401	115.623	-6.517	1.00	33.98	T5
	ATOM	4862	N	GLU	A	38	28.303	115.806	-4.276	1.00	35.35	T5
	ATOM	4863	CA	GLU	A	38	29.681	116.277	-4.110	1.00	29.47	T5
	ATOM	4864	CB	GLU	A	38	29.709	117.800	-4.063	1.00	37.16	T5
65	ATOM	4865	CG	GLU	A	38	29.676	118.499	-5.404	1.00	35.52	T5
	ATOM	4866	CD	GLU	A	38	29.451	120.003	-5.262	1.00	33.49	T5

	ATOM	4867	OE1	GLU	A	38	29.960	120.600	-4.275	1.00	32.50	T5
	ATOM	4868	OE2	GLU	A	38	28.772	120.588	-6.144	1.00	29.69	T5
	ATOM	4869	C	GLU	A	38	30.317	115.774	-2.826	1.00	34.89	T5
	ATOM	4870	O	GLU	A	38	29.618	115.311	-1.927	1.00	31.88	T5
5	ATOM	4871	N	GLU	A	39	31.643	115.868	-2.741	1.00	28.54	T5
	ATOM	4872	CA	GLU	A	39	32.339	115.466	-1.524	1.00	28.61	T5
	ATOM	4873	CB	GLU	A	39	33.761	115.017	-1.783	1.00	26.51	T5
	ATOM	4874	CG	GLU	A	39	33.969	114.142	-2.955	1.00	32.10	T5
	ATOM	4875	CD	GLU	A	39	35.361	114.337	-3.525	1.00	28.58	T5
10	ATOM	4876	OE1	GLU	A	39	35.498	115.113	-4.508	1.00	33.83	T5
	ATOM	4877	OE2	GLU	A	39	36.316	113.734	-2.972	1.00	35.68	T5
	ATOM	4878	C	GLU	A	39	32.463	116.727	-0.703	1.00	30.61	T5
	ATOM	4879	O	GLU	A	39	32.640	117.812	-1.251	1.00	31.17	T5
	ATOM	4880	N	LYS	A	40	32.393	116.596	0.609	1.00	34.63	T5
15	ATOM	4881	CA	LYS	A	40	32.543	117.761	1.448	1.00	32.00	T5
	ATOM	4882	CB	LYS	A	40	31.257	118.583	1.478	1.00	27.72	T5
	ATOM	4883	CG	LYS	A	40	31.394	119.853	2.316	1.00	27.75	T5
	ATOM	4884	CD	LYS	A	40	30.042	120.442	2.660	1.00	32.04	T5
	ATOM	4885	CE	LYS	A	40	30.170	121.576	3.673	1.00	27.22	T5
20	ATOM	4886	NZ	LYS	A	40	28.823	122.051	4.132	1.00	31.39	T5
	ATOM	4887	C	LYS	A	40	32.919	117.338	2.852	1.00	30.88	T5
	ATOM	4888	O	LYS	A	40	32.085	116.848	3.612	1.00	26.46	T5
	ATOM	4889	N	GLU	A	41	34.192	117.512	3.182	1.00	29.02	T5
	ATOM	4890	CA	GLU	A	41	34.681	117.171	4.501	1.00	27.60	T5
25	ATOM	4891	CB	GLU	A	41	34.171	118.208	5.492	1.00	25.52	T5
	ATOM	4892	CG	GLU	A	41	34.458	119.623	5.014	1.00	37.74	T5
	ATOM	4893	CD	GLU	A	41	33.673	120.678	5.769	1.00	29.90	T5
	ATOM	4894	OE1	GLU	A	41	32.425	120.545	5.875	1.00	33.33	T5
	ATOM	4895	OE2	GLU	A	41	34.307	121.649	6.249	1.00	24.59	T5
30	ATOM	4896	C	GLU	A	41	34.268	115.766	4.916	1.00	29.69	T5
	ATOM	4897	O	GLU	A	41	33.590	115.571	5.921	1.00	30.37	T5
	ATOM	4898	N	ASN	A	42	34.678	114.794	4.111	1.00	33.98	T5
	ATOM	4899	CA	ASN	A	42	34.405	113.394	4.374	1.00	32.18	T5
	ATOM	4900	CB	ASN	A	42	35.045	112.988	5.685	1.00	26.61	T5
35	ATOM	4901	CG	ASN	A	42	35.709	111.656	5.596	1.00	30.11	T5
	ATOM	4902	OD1	ASN	A	42	35.446	110.769	6.403	1.00	26.91	T5
	ATOM	4903	ND2	ASN	A	42	36.587	111.496	4.608	1.00	33.26	T5
	ATOM	4904	C	ASN	A	42	32.943	113.000	4.398	1.00	34.19	T5
	ATOM	4905	O	ASN	A	42	32.591	111.965	4.949	1.00	30.54	T5
40	ATOM	4906	N	LYS	A	43	32.096	113.824	3.797	1.00	26.86	T5
	ATOM	4907	CA	LYS	A	43	30.668	113.547	3.732	1.00	24.51	T5
	ATOM	4908	CB	LYS	A	43	29.908	114.465	4.684	1.00	30.37	T5
	ATOM	4909	CG	LYS	A	43	30.149	114.177	6.144	1.00	27.62	T5
	ATOM	4910	CD	LYS	A	43	29.418	115.165	7.034	1.00	33.25	T5
45	ATOM	4911	CE	LYS	A	43	30.067	116.532	6.961	1.00	31.67	T5
	ATOM	4912	NZ	LYS	A	43	29.414	117.510	7.865	1.00	29.02	T5
	ATOM	4913	C	LYS	A	43	30.185	113.783	2.311	1.00	30.26	T5
	ATOM	4914	O	LYS	A	43	30.852	114.448	1.523	1.00	34.47	T5
	ATOM	4915	N	ILE	A	44	29.030	113.230	1.978	1.00	34.65	T5
50	ATOM	4916	CA	ILE	A	44	28.480	113.429	0.652	1.00	37.50	T5
	ATOM	4917	CB	ILE	A	44	27.799	112.170	0.136	1.00	33.73	T5
	ATOM	4918	CG2	ILE	A	44	27.239	112.422	-1.243	1.00	35.54	T5
	ATOM	4919	CG1	ILE	A	44	28.811	111.030	0.079	1.00	27.81	T5
	ATOM	4920	CD1	ILE	A	44	28.239	109.725	-0.392	1.00	29.33	T5
55	ATOM	4921	C	ILE	A	44	27.459	114.549	0.744	1.00	26.71	T5
	ATOM	4922	O	ILE	A	44	26.527	114.484	1.540	1.00	29.85	T5
	ATOM	4923	N	LEU	A	45	27.646	115.580	-0.069	1.00	25.10	T5
	ATOM	4924	CA	LEU	A	45	26.753	116.724	-0.069	1.00	36.19	T5
	ATOM	4925	CB	LEU	A	45	27.570	118.002	-0.192	1.00	25.88	T5
60	ATOM	4926	CG	LEU	A	45	26.727	119.271	-0.277	1.00	24.29	T5
	ATOM	4927	CD1	LEU	A	45	26.072	119.550	1.079	1.00	36.67	T5
	ATOM	4928	CD2	LEU	A	45	27.602	120.424	-0.700	1.00	37.35	T5
	ATOM	4929	C	LEU	A	45	25.715	116.679	-1.189	1.00	31.60	T5
	ATOM	4930	O	LEU	A	45	26.056	116.501	-2.354	1.00	31.45	T5
65	ATOM	4931	N	VAL	A	46	24.447	116.855	-0.830	1.00	29.73	T5
	ATOM	4932	CA	VAL	A	46	23.369	116.832	-1.805	1.00	33.32	T5

	ATOM	4933	CB	VAL	A	46	22.051	116.410	-1.148	1.00	29.29	T5
	ATOM	4934	CG1	VAL	A	46	20.941	116.407	-2.170	1.00	24.32	T5
	ATOM	4935	CG2	VAL	A	46	22.199	115.042	-0.538	1.00	35.88	T5
	ATOM	4936	C	VAL	A	46	23.186	118.206	-2.441	1.00	30.98	T5
5	ATOM	4937	O	VAL	A	46	22.901	119.184	-1.755	1.00	30.21	T5
	ATOM	4938	N	LYS	A	47	23.341	118.277	-3.758	1.00	29.60	T5
	ATOM	4939	CA	LYS	A	47	23.198	119.545	-4.463	1.00	24.68	T5
	ATOM	4940	CB	LYS	A	47	24.342	119.710	-5.465	1.00	30.96	T5
	ATOM	4941	CG	LYS	A	47	25.668	120.085	-4.842	1.00	28.69	T5
10	ATOM	4942	CD	LYS	A	47	25.502	121.315	-3.973	1.00	33.05	T5
	ATOM	4943	CE	LYS	A	47	26.832	121.926	-3.571	1.00	32.94	T5
	ATOM	4944	NZ	LYS	A	47	27.517	122.619	-4.715	1.00	31.19	T5
	ATOM	4945	C	LYS	A	47	21.861	119.744	-5.179	1.00	32.60	T5
	ATOM	4946	O	LYS	A	47	21.571	120.829	-5.661	1.00	25.38	T5
15	ATOM	4947	N	GLU	A	48	21.055	118.695	-5.254	1.00	34.47	T5
	ATOM	4948	CA	GLU	A	48	19.759	118.762	-5.912	1.00	26.10	T5
	ATOM	4949	CB	GLU	A	48	19.837	118.226	-7.328	1.00	31.53	T5
	ATOM	4950	CG	GLU	A	48	20.586	119.083	-8.300	1.00	35.25	T5
	ATOM	4951	CD	GLU	A	48	20.735	118.393	-9.652	1.00	26.86	T5
20	ATOM	4952	OE1	GLU	A	48	19.743	117.764	-10.109	1.00	25.86	T5
	ATOM	4953	OE2	GLU	A	48	21.839	118.481	-10.255	1.00	34.30	T5
	ATOM	4954	C	GLU	A	48	18.834	117.861	-5.148	1.00	35.28	T5
	ATOM	4955	O	GLU	A	48	19.139	116.679	-4.974	1.00	28.24	T5
	ATOM	4956	N	THR	A	49	17.700	118.386	-4.701	1.00	29.29	T5
25	ATOM	4957	CA	THR	A	49	16.786	117.537	-3.959	1.00	28.77	T5
	ATOM	4958	CB	THR	A	49	15.680	118.355	-3.300	1.00	37.96	T5
	ATOM	4959	OG1	THR	A	49	14.571	118.434	-4.190	1.00	26.93	T5
	ATOM	4960	CG2	THR	A	49	16.174	119.759	-2.978	1.00	30.34	T5
	ATOM	4961	C	THR	A	49	16.174	116.490	-4.897	1.00	32.17	T5
30	ATOM	4962	O	THR	A	49	16.018	116.721	-6.094	1.00	30.13	T5
	ATOM	4963	N	GLY	A	50	15.857	115.326	-4.345	1.00	24.37	T5
	ATOM	4964	CA	GLY	A	50	15.272	114.262	-5.137	1.00	30.00	T5
	ATOM	4965	C	GLY	A	50	15.358	112.945	-4.398	1.00	35.59	T5
	ATOM	4966	O	GLY	A	50	15.603	112.927	-3.191	1.00	31.96	T5
35	ATOM	4967	N	TYR	A	51	15.155	111.844	-5.114	1.00	34.92	T5
	ATOM	4968	CA	TYR	A	51	15.229	110.517	-4.511	1.00	25.96	T5
	ATOM	4969	CB	TYR	A	51	14.073	109.637	-5.002	1.00	32.20	T5
	ATOM	4970	CG	TYR	A	51	12.760	110.014	-4.375	1.00	27.31	T5
	ATOM	4971	CD1	TYR	A	51	12.037	111.116	-4.829	1.00	20.80	T5
40	ATOM	4972	CE1	TYR	A	51	10.876	111.544	-4.170	1.00	24.45	T5
	ATOM	4973	CD2	TYR	A	51	12.289	109.333	-3.252	1.00	32.34	T5
	ATOM	4974	CE2	TYR	A	51	11.138	109.746	-2.586	1.00	36.95	T5
	ATOM	4975	CZ	TYR	A	51	10.436	110.857	-3.044	1.00	33.18	T5
	ATOM	4976	OH	TYR	A	51	9.333	111.308	-2.340	1.00	26.86	T5
45	ATOM	4977	C	TYR	A	51	16.566	109.856	-4.829	1.00	34.25	T5
	ATOM	4978	O	TYR	A	51	16.980	109.785	-5.993	1.00	33.54	T5
	ATOM	4979	N	PHE	A	52	17.240	109.371	-3.792	1.00	29.59	T5
	ATOM	4980	CA	PHE	A	52	18.538	108.739	-3.980	1.00	35.99	T5
	ATOM	4981	CB	PHE	A	52	19.643	109.582	-3.334	1.00	31.37	T5
50	ATOM	4982	CG	PHE	A	52	19.731	110.986	-3.852	1.00	24.34	T5
	ATOM	4983	CD1	PHE	A	52	18.806	111.940	-3.463	1.00	29.53	T5
	ATOM	4984	CD2	PHE	A	52	20.749	111.355	-4.715	1.00	37.58	T5
	ATOM	4985	CE1	PHE	A	52	18.895	113.249	-3.926	1.00	32.06	T5
	ATOM	4986	CE2	PHE	A	52	20.845	112.655	-5.180	1.00	33.80	T5
55	ATOM	4987	CZ	PHE	A	52	19.916	113.606	-4.785	1.00	38.41	T5
	ATOM	4988	C	PHE	A	52	18.639	107.331	-3.417	1.00	25.47	T5
	ATOM	4989	O	PHE	A	52	18.005	107.001	-2.417	1.00	25.12	T5
	ATOM	4990	N	PHE	A	53	19.440	106.511	-4.089	1.00	38.63	T5
	ATOM	4991	CA	PHE	A	53	19.713	105.150	-3.662	1.00	33.80	T5
60	ATOM	4992	CB	PHE	A	53	19.889	104.232	-4.861	1.00	30.20	T5
	ATOM	4993	CG	PHE	A	53	20.380	102.863	-4.502	1.00	29.01	T5
	ATOM	4994	CD1	PHE	A	53	19.579	101.988	-3.781	1.00	27.25	T5
	ATOM	4995	CD2	PHE	A	53	21.655	102.449	-4.869	1.00	26.95	T5
	ATOM	4996	CE1	PHE	A	53	20.044	100.722	-3.432	1.00	31.98	T5
65	ATOM	4997	CE2	PHE	A	53	22.123	101.189	-4.524	1.00	33.30	T5
	ATOM	4998	CZ	PHE	A	53	21.316	100.327	-3.806	1.00	26.27	T5

	ATOM	4999	C	PHE	A	53	21.044	105.305	-2.941	1.00	29.74	T5
	ATOM	5000	O	PHE	A	53	22.000	105.799	-3.517	1.00	27.04	T5
	ATOM	5001	N	ILE	A	54	21.112	104.893	-1.685	1.00	27.53	T5
	ATOM	5002	CA	ILE	A	54	22.337	105.052	-0.924	1.00	25.14	T5
5	ATOM	5003	CB	ILE	A	54	22.091	105.982	0.265	1.00	31.93	T5
	ATOM	5004	CG2	ILE	A	54	23.395	106.314	0.946	1.00	34.18	T5
	ATOM	5005	CG1	ILE	A	54	21.422	107.261	-0.231	1.00	24.10	T5
	ATOM	5006	CD1	ILE	A	54	20.722	108.033	0.845	1.00	23.57	T5
	ATOM	5007	C	ILE	A	54	22.841	103.712	-0.427	1.00	28.96	T5
10	ATOM	5008	O	ILE	A	54	22.083	102.933	0.142	1.00	31.61	T5
	ATOM	5009	N	TYR	A	55	24.123	103.442	-0.641	1.00	35.48	T5
	ATOM	5010	CA	TYR	A	55	24.701	102.176	-0.215	1.00	30.04	T5
	ATOM	5011	CB	TYR	A	55	24.924	101.281	-1.430	1.00	28.02	T5
	ATOM	5012	CG	TYR	A	55	25.783	101.902	-2.498	1.00	27.39	T5
15	ATOM	5013	CD1	TYR	A	55	27.141	101.642	-2.558	1.00	31.50	T5
	ATOM	5014	CE1	TYR	A	55	27.944	102.234	-3.523	1.00	27.30	T5
	ATOM	5015	CD2	TYR	A	55	25.243	102.770	-3.430	1.00	28.94	T5
	ATOM	5016	CE2	TYR	A	55	26.034	103.368	-4.396	1.00	30.54	T5
	ATOM	5017	CZ	TYR	A	55	27.384	103.099	-4.437	1.00	33.14	T5
20	ATOM	5018	OH	TYR	A	55	28.179	103.710	-5.376	1.00	35.99	T5
	ATOM	5019	C	TYR	A	55	26.006	102.360	0.541	1.00	29.59	T5
	ATOM	5020	O	TYR	A	55	26.535	103.456	0.621	1.00	32.76	T5
	ATOM	5021	N	GLY	A	56	26.519	101.277	1.105	1.00	28.46	T5
	ATOM	5022	CA	GLY	A	56	27.759	101.360	1.841	1.00	31.45	T5
25	ATOM	5023	C	GLY	A	56	28.132	100.034	2.456	1.00	32.06	T5
	ATOM	5024	O	GLY	A	56	27.272	99.321	2.955	1.00	31.95	T5
	ATOM	5025	N	GLN	A	57	29.420	99.707	2.398	1.00	31.93	T5
	ATOM	5026	CA	GLN	A	57	29.949	98.470	2.963	1.00	31.07	T5
	ATOM	5027	CB	GLN	A	57	30.272	97.459	1.859	1.00	36.80	T5
30	ATOM	5028	CG	GLN	A	57	30.977	96.211	2.369	1.00	31.56	T5
	ATOM	5029	CD	GLN	A	57	31.183	95.153	1.303	1.00	28.30	T5
	ATOM	5030	OE1	GLN	A	57	30.233	94.520	0.846	1.00	28.56	T5
	ATOM	5031	NE2	GLN	A	57	32.430	94.953	0.900	1.00	30.92	T5
	ATOM	5032	C	GLN	A	57	31.217	98.758	3.768	1.00	23.78	T5
35	ATOM	5033	O	GLN	A	57	31.980	99.675	3.451	1.00	31.30	T5
	ATOM	5034	N	VAL	A	58	31.428	97.971	4.817	1.00	24.85	T5
	ATOM	5035	CA	VAL	A	58	32.594	98.106	5.682	1.00	32.42	T5
	ATOM	5036	CB	VAL	A	58	32.238	98.858	6.989	1.00	26.52	T5
	ATOM	5037	CG1	VAL	A	58	33.365	98.744	7.988	1.00	32.83	T5
40	ATOM	5038	CG2	VAL	A	58	31.965	100.305	6.687	1.00	32.66	T5
	ATOM	5039	C	VAL	A	58	33.072	96.711	6.053	1.00	28.35	T5
	ATOM	5040	O	VAL	A	58	32.255	95.832	6.300	1.00	26.50	T5
	ATOM	5041	N	LEU	A	59	34.388	96.503	6.082	1.00	25.15	T5
	ATOM	5042	CA	LEU	A	59	34.947	95.204	6.466	1.00	27.75	T5
45	ATOM	5043	CB	LEU	A	59	36.142	94.849	5.586	1.00	27.87	T5
	ATOM	5044	CG	LEU	A	59	36.468	93.360	5.449	1.00	33.93	T5
	ATOM	5045	CD1	LEU	A	59	37.877	93.211	4.915	1.00	27.12	T5
	ATOM	5046	CD2	LEU	A	59	36.345	92.654	6.783	1.00	28.58	T5
	ATOM	5047	C	LEU	A	59	35.402	95.292	7.927	1.00	30.10	T5
50	ATOM	5048	O	LEU	A	59	36.329	96.040	8.251	1.00	27.45	T5
	ATOM	5049	N	TYR	A	60	34.751	94.535	8.806	1.00	32.41	T5
	ATOM	5050	CA	TYR	A	60	35.108	94.558	10.217	1.00	30.66	T5
	ATOM	5051	CB	TYR	A	60	33.862	94.389	11.073	1.00	38.39	T5
	ATOM	5052	CG	TYR	A	60	32.873	95.481	10.839	1.00	31.46	T5
55	ATOM	5053	CD1	TYR	A	60	31.716	95.249	10.108	1.00	28.05	T5
	ATOM	5054	CE1	TYR	A	60	30.831	96.284	9.816	1.00	27.45	T5
	ATOM	5055	CD2	TYR	A	60	33.130	96.772	11.284	1.00	26.72	T5
	ATOM	5056	CE2	TYR	A	60	32.268	97.812	11.003	1.00	36.90	T5
	ATOM	5057	CZ	TYR	A	60	31.120	97.567	10.266	1.00	21.75	T5
60	ATOM	5058	OH	TYR	A	60	30.273	98.614	9.970	1.00	26.98	T5
	ATOM	5059	C	TYR	A	60	36.124	93.501	10.606	1.00	26.12	T5
	ATOM	5060	O	TYR	A	60	35.936	92.315	10.343	1.00	35.88	T5
	ATOM	5061	N	THR	A	61	37.204	93.941	11.240	1.00	33.21	T5
	ATOM	5062	CA	THR	A	61	38.249	93.034	11.682	1.00	34.43	T5
65	ATOM	5063	CB	THR	A	61	39.578	93.352	11.007	1.00	32.30	T5
	ATOM	5064	OG1	THR	A	61	39.902	94.725	11.220	1.00	25.25	T5

	ATOM	5065	CG2	THR	A	61	39.488	93.087	9.520	1.00	34.90	T5
	ATOM	5066	C	THR	A	61	38.391	93.178	13.186	1.00	31.90	T5
	ATOM	5067	O	THR	A	61	39.374	92.762	13.778	1.00	34.37	T5
	ATOM	5068	N	ASP	A	62	37.379	93.778	13.793	1.00	32.92	T5
5	ATOM	5069	CA	ASP	A	62	37.329	93.988	15.231	1.00	27.67	T5
	ATOM	5070	CB	ASP	A	62	36.462	95.215	15.514	1.00	36.40	T5
	ATOM	5071	CG	ASP	A	62	36.590	95.711	16.932	1.00	30.94	T5
	ATOM	5072	OD1	ASP	A	62	36.816	96.937	17.104	1.00	38.82	T5
	ATOM	5073	OD2	ASP	A	62	36.455	94.879	17.863	1.00	33.61	T5
10	ATOM	5074	C	ASP	A	62	36.702	92.734	15.839	1.00	22.81	T5
	ATOM	5075	O	ASP	A	62	35.866	92.091	15.206	1.00	35.62	T5
	ATOM	5076	N	LYS	A	63	37.088	92.369	17.056	1.00	26.10	T5
	ATOM	5077	CA	LYS	A	63	36.514	91.163	17.651	1.00	29.38	T5
	ATOM	5078	CB	LYS	A	63	37.623	90.243	18.160	1.00	30.20	T5
15	ATOM	5079	CG	LYS	A	63	38.506	90.878	19.216	1.00	33.19	T5
	ATOM	5080	CD	LYS	A	63	39.574	89.898	19.713	1.00	31.22	T5
	ATOM	5081	CE	LYS	A	63	40.520	89.448	18.588	1.00	27.45	T5
	ATOM	5082	NZ	LYS	A	63	41.560	88.465	19.042	1.00	32.41	T5
	ATOM	5083	C	LYS	A	63	35.529	91.421	18.780	1.00	28.30	T5
20	ATOM	5084	O	LYS	A	63	35.290	90.541	19.610	1.00	30.20	T5
	ATOM	5085	N	THR	A	64	34.930	92.606	18.811	1.00	30.98	T5
	ATOM	5086	CA	THR	A	64	33.998	92.895	19.893	1.00	29.96	T5
	ATOM	5087	CB	THR	A	64	33.994	94.411	20.267	1.00	43.68	T5
	ATOM	5088	OG1	THR	A	64	33.791	95.205	19.095	1.00	28.42	T5
25	ATOM	5089	CG2	THR	A	64	35.316	94.796	20.930	1.00	29.32	T5
	ATOM	5090	C	THR	A	64	32.558	92.413	19.710	1.00	24.26	T5
	ATOM	5091	O	THR	A	64	31.623	93.206	19.714	1.00	26.31	T5
	ATOM	5092	N	TYR	A	65	32.403	91.100	19.564	1.00	30.39	T5
	ATOM	5093	CA	TYR	A	65	31.098	90.437	19.433	1.00	32.52	T5
30	ATOM	5094	CB	TYR	A	65	30.473	90.279	20.828	1.00	26.32	T5
	ATOM	5095	CG	TYR	A	65	29.375	91.273	21.147	1.00	31.04	T5
	ATOM	5096	CD1	TYR	A	65	28.032	90.971	20.893	1.00	34.79	T5
	ATOM	5097	CE1	TYR	A	65	27.010	91.888	21.188	1.00	27.98	T5
	ATOM	5098	CD2	TYR	A	65	29.677	92.516	21.700	1.00	32.81	T5
35	ATOM	5099	CE2	TYR	A	65	28.670	93.445	21.997	1.00	32.46	T5
	ATOM	5100	CZ	TYR	A	65	27.339	93.128	21.741	1.00	35.26	T5
	ATOM	5101	OH	TYR	A	65	26.350	94.052	22.042	1.00	35.95	T5
	ATOM	5102	C	TYR	A	65	30.041	91.011	18.476	1.00	29.82	T5
	ATOM	5103	O	TYR	A	65	29.165	90.277	18.017	1.00	29.77	T5
40	ATOM	5104	N	ALA	A	66	30.101	92.303	18.179	1.00	30.02	T5
	ATOM	5105	CA	ALA	A	66	29.117	92.893	17.278	1.00	33.48	T5
	ATOM	5106	CB	ALA	A	66	27.803	93.080	18.009	1.00	29.63	T5
	ATOM	5107	C	ALA	A	66	29.590	94.220	16.712	1.00	32.67	T5
	ATOM	5108	O	ALA	A	66	29.830	95.170	17.452	1.00	30.18	T5
45	ATOM	5109	N	MET	A	67	29.731	94.278	15.395	1.00	31.50	T5
	ATOM	5110	CA	MET	A	67	30.163	95.501	14.740	1.00	28.26	T5
	ATOM	5111	CB	MET	A	67	31.490	95.277	14.017	1.00	30.85	T5
	ATOM	5112	CG	MET	A	67	32.664	95.088	14.951	1.00	23.23	T5
	ATOM	5113	SD	MET	A	67	32.878	96.524	16.027	1.00	33.81	T5
50	ATOM	5114	CE	MET	A	67	33.755	97.643	14.914	1.00	28.68	T5
	ATOM	5115	C	MET	A	67	29.097	95.927	13.749	1.00	31.83	T5
	ATOM	5116	O	MET	A	67	28.183	95.162	13.462	1.00	35.24	T5
	ATOM	5117	N	GLY	A	68	29.212	97.147	13.235	1.00	28.47	T5
	ATOM	5118	CA	GLY	A	68	28.245	97.633	12.272	1.00	30.54	T5
55	ATOM	5119	C	GLY	A	68	28.398	99.114	12.006	1.00	28.36	T5
	ATOM	5120	O	GLY	A	68	29.152	99.796	12.702	1.00	27.55	T5
	ATOM	5121	N	HIS	A	69	27.702	99.618	10.989	1.00	26.34	T5
	ATOM	5122	CA	HIS	A	69	27.761	101.040	10.675	1.00	31.98	T5
	ATOM	5123	CB	HIS	A	69	28.695	101.307	9.492	1.00	31.73	T5
60	ATOM	5124	CG	HIS	A	69	28.374	100.515	8.267	1.00	22.55	T5
	ATOM	5125	CD2	HIS	A	69	27.790	100.875	7.101	1.00	29.98	T5
	ATOM	5126	ND1	HIS	A	69	28.701	99.185	8.135	1.00	30.77	T5
	ATOM	5127	CE1	HIS	A	69	28.339	98.760	6.938	1.00	32.42	T5
	ATOM	5128	NE2	HIS	A	69	27.783	99.767	6.290	1.00	41.24	T5
65	ATOM	5129	C	HIS	A	69	26.390	101.649	10.400	1.00	26.70	T5
	ATOM	5130	O	HIS	A	69	25.396	100.940	10.246	1.00	32.48	T5

	ATOM	5131	N	LEU	A	70	26.352	102.975	10.350	1.00	37.99	T5
	ATOM	5132	CA	LEU	A	70	25.121	103.706	10.112	1.00	30.41	T5
	ATOM	5133	CB	LEU	A	70	24.777	104.567	11.325	1.00	37.35	T5
	ATOM	5134	CG	LEU	A	70	24.995	103.992	12.714	1.00	36.47	T5
5	ATOM	5135	CD1	LEU	A	70	24.778	105.087	13.722	1.00	27.85	T5
	ATOM	5136	CD2	LEU	A	70	24.056	102.843	12.962	1.00	35.04	T5
	ATOM	5137	C	LEU	A	70	25.278	104.640	8.933	1.00	25.27	T5
	ATOM	5138	O	LEU	A	70	26.349	105.206	8.720	1.00	32.86	T5
	ATOM	5139	N	ILE	A	71	24.214	104.802	8.162	1.00	29.17	T5
10	ATOM	5140	CA	ILE	A	71	24.239	105.737	7.052	1.00	30.92	T5
	ATOM	5141	CB	ILE	A	71	23.714	105.103	5.763	1.00	36.80	T5
	ATOM	5142	CG2	ILE	A	71	23.461	106.171	4.730	1.00	28.74	T5
	ATOM	5143	CG1	ILE	A	71	24.737	104.098	5.235	1.00	30.33	T5
	ATOM	5144	CD1	ILE	A	71	24.308	103.387	3.974	1.00	30.32	T5
15	ATOM	5145	C	ILE	A	71	23.308	106.843	7.533	1.00	22.84	T5
	ATOM	5146	O	ILE	A	71	22.101	106.645	7.635	1.00	30.99	T5
	ATOM	5147	N	GLN	A	72	23.869	108.005	7.846	1.00	34.81	T5
	ATOM	5148	CA	GLN	A	72	23.061	109.096	8.364	1.00	29.95	T5
	ATOM	5149	CB	GLN	A	72	23.653	109.557	9.682	1.00	27.39	T5
20	ATOM	5150	CG	GLN	A	72	24.014	108.392	10.579	1.00	25.75	T5
	ATOM	5151	CD	GLN	A	72	24.412	108.837	11.965	1.00	24.82	T5
	ATOM	5152	OE1	GLN	A	72	25.315	109.660	12.127	1.00	33.34	T5
	ATOM	5153	NE2	GLN	A	72	23.742	108.296	12.979	1.00	29.15	T5
	ATOM	5154	C	GLN	A	72	22.844	110.290	7.455	1.00	29.21	T5
25	ATOM	5155	O	GLN	A	72	23.612	110.545	6.538	1.00	30.04	T5
	ATOM	5156	N	ARG	A	73	21.775	111.020	7.744	1.00	29.71	T5
	ATOM	5157	CA	ARG	A	73	21.393	112.197	6.988	1.00	29.64	T5
	ATOM	5158	CB	ARG	A	73	20.046	111.945	6.327	1.00	33.75	T5
	ATOM	5159	CG	ARG	A	73	19.537	113.100	5.528	1.00	31.50	T5
30	ATOM	5160	CD	ARG	A	73	18.041	113.050	5.408	1.00	32.89	T5
	ATOM	5161	NE	ARG	A	73	17.537	114.190	4.655	1.00	32.05	T5
	ATOM	5162	CZ	ARG	A	73	16.265	114.560	4.636	1.00	28.12	T5
	ATOM	5163	NH1	ARG	A	73	15.364	113.880	5.332	1.00	29.08	T5
	ATOM	5164	NH2	ARG	A	73	15.900	115.606	3.921	1.00	26.52	T5
35	ATOM	5165	C	ARG	A	73	21.290	113.429	7.898	1.00	27.52	T5
	ATOM	5166	O	ARG	A	73	20.693	113.367	8.974	1.00	34.70	T5
	ATOM	5167	N	LYS	A	74	21.887	114.539	7.466	1.00	23.18	T5
	ATOM	5168	CA	LYS	A	74	21.832	115.786	8.222	1.00	31.34	T5
	ATOM	5169	CB	LYS	A	74	23.210	116.437	8.309	1.00	29.80	T5
40	ATOM	5170	CG	LYS	A	74	24.212	115.673	9.160	1.00	27.16	T5
	ATOM	5171	CD	LYS	A	74	25.611	116.324	9.147	1.00	32.84	T5
	ATOM	5172	CE	LYS	A	74	26.622	115.511	9.982	1.00	34.42	T5
	ATOM	5173	NZ	LYS	A	74	27.993	116.123	10.020	1.00	26.32	T5
	ATOM	5174	C	LYS	A	74	20.893	116.735	7.504	1.00	34.53	T5
45	ATOM	5175	O	LYS	A	74	21.272	117.358	6.509	1.00	24.25	T5
	ATOM	5176	N	LYS	A	75	19.669	116.839	8.006	1.00	32.05	T5
	ATOM	5177	CA	LYS	A	75	18.673	117.712	7.407	1.00	25.59	T5
	ATOM	5178	CB	LYS	A	75	17.360	117.623	8.185	1.00	30.13	T5
	ATOM	5179	CG	LYS	A	75	16.715	116.256	8.239	1.00	25.51	T5
50	ATOM	5180	CD	LYS	A	75	15.414	116.343	9.029	1.00	26.27	T5
	ATOM	5181	CE	LYS	A	75	14.735	114.981	9.125	1.00	25.29	T5
	ATOM	5182	NZ	LYS	A	75	13.510	114.958	9.999	1.00	29.52	T5
	ATOM	5183	C	LYS	A	75	19.123	119.174	7.389	1.00	28.86	T5
	ATOM	5184	O	LYS	A	75	19.648	119.669	8.380	1.00	30.25	T5
55	ATOM	5185	N	VAL	A	76	18.905	119.864	6.271	1.00	31.13	T5
	ATOM	5186	CA	VAL	A	76	19.270	121.275	6.166	1.00	30.82	T5
	ATOM	5187	CB	VAL	A	76	19.257	121.788	4.734	1.00	24.29	T5
	ATOM	5188	CG1	VAL	A	76	20.009	123.078	4.663	1.00	29.92	T5
	ATOM	5189	CG2	VAL	A	76	19.822	120.772	3.810	1.00	27.78	T5
60	ATOM	5190	C	VAL	A	76	18.185	122.057	6.869	1.00	24.62	T5
	ATOM	5191	O	VAL	A	76	18.458	123.029	7.577	1.00	35.85	T5
	ATOM	5192	N	HIS	A	77	16.946	121.627	6.644	1.00	28.34	T5
	ATOM	5193	CA	HIS	A	77	15.787	122.267	7.233	1.00	29.76	T5
	ATOM	5194	CB	HIS	A	77	14.711	122.464	6.165	1.00	41.41	T5
65	ATOM	5195	CG	HIS	A	77	15.166	123.282	4.997	1.00	28.24	T5
	ATOM	5196	CD2	HIS	A	77	14.689	123.377	3.735	1.00	29.10	T5

	ATOM	5197	ND1	HIS	A	77	16.205	124.183	5.084	1.00	29.04	T5
	ATOM	5198	CE1	HIS	A	77	16.348	124.801	3.926	1.00	33.18	T5
	ATOM	5199	NE2	HIS	A	77	15.439	124.331	3.092	1.00	27.79	T5
	ATOM	5200	C	HIS	A	77	15.275	121.401	8.372	1.00	37.60	T5
5	ATOM	5201	O	HIS	A	77	15.367	120.177	8.307	1.00	29.87	T5
	ATOM	5202	N	VAL	A	78	14.721	122.027	9.407	1.00	28.62	T5
	ATOM	5203	CA	VAL	A	78	14.257	121.254	10.544	1.00	29.10	T5
	ATOM	5204	CB	VAL	A	78	15.212	121.479	11.734	1.00	30.08	T5
	ATOM	5205	CG1	VAL	A	78	14.852	120.572	12.881	1.00	28.94	T5
10	ATOM	5206	CG2	VAL	A	78	16.635	121.169	11.307	1.00	28.92	T5
	ATOM	5207	C	VAL	A	78	12.788	121.349	11.015	1.00	36.30	T5
	ATOM	5208	O	VAL	A	78	12.058	120.366	10.926	1.00	33.11	T5
	ATOM	5209	N	PHE	A	79	12.334	122.485	11.520	1.00	28.17	T5
	ATOM	5210	CA	PHE	A	79	10.939	122.580	11.983	1.00	28.56	T5
15	ATOM	5211	CB	PHE	A	79	9.957	121.919	11.006	1.00	27.54	T5
	ATOM	5212	CG	PHE	A	79	10.072	122.397	9.598	1.00	26.56	T5
	ATOM	5213	CD1	PHE	A	79	10.733	121.632	8.651	1.00	26.29	T5
	ATOM	5214	CD2	PHE	A	79	9.525	123.618	9.216	1.00	33.89	T5
	ATOM	5215	CE1	PHE	A	79	10.853	122.067	7.350	1.00	32.35	T5
20	ATOM	5216	CE2	PHE	A	79	9.638	124.068	7.911	1.00	24.08	T5
	ATOM	5217	CZ	PHE	A	79	10.304	123.291	6.977	1.00	29.84	T5
	ATOM	5218	C	PHE	A	79	10.682	121.939	13.350	1.00	28.02	T5
	ATOM	5219	O	PHE	A	79	10.930	120.751	13.551	1.00	35.58	T5
	ATOM	5220	N	GLY	A	80	10.150	122.737	14.273	1.00	32.55	T5
25	ATOM	5221	CA	GLY	A	80	9.820	122.255	15.606	1.00	27.94	T5
	ATOM	5222	C	GLY	A	80	10.854	121.418	16.329	1.00	29.71	T5
	ATOM	5223	O	GLY	A	80	12.021	121.785	16.431	1.00	29.34	T5
	ATOM	5224	N	ASP	A	81	10.411	120.277	16.841	1.00	22.92	T5
	ATOM	5225	CA	ASP	A	81	11.280	119.371	17.579	1.00	27.63	T5
30	ATOM	5226	CB	ASP	A	81	10.541	118.823	18.805	1.00	26.45	T5
	ATOM	5227	CG	ASP	A	81	9.359	117.937	18.445	1.00	22.81	T5
	ATOM	5228	OD1	ASP	A	81	8.904	117.974	17.290	1.00	31.96	T5
	ATOM	5229	OD2	ASP	A	81	8.866	117.206	19.327	1.00	31.27	T5
	ATOM	5230	C	ASP	A	81	11.843	118.212	16.766	1.00	35.47	T5
35	ATOM	5231	O	ASP	A	81	12.107	117.141	17.314	1.00	27.55	T5
	ATOM	5232	N	GLU	A	82	12.021	118.419	15.463	1.00	32.59	T5
	ATOM	5233	CA	GLU	A	82	12.583	117.377	14.610	1.00	34.42	T5
	ATOM	5234	CB	GLU	A	82	12.538	117.770	13.139	1.00	37.04	T5
	ATOM	5235	CG	GLU	A	82	11.242	117.598	12.426	1.00	30.32	T5
40	ATOM	5236	CD	GLU	A	82	11.480	117.361	10.951	1.00	35.81	T5
	ATOM	5237	OE1	GLU	A	82	12.343	118.056	10.378	1.00	31.76	T5
	ATOM	5238	OE2	GLU	A	82	10.815	116.481	10.362	1.00	35.65	T5
	ATOM	5239	C	GLU	A	82	14.047	117.250	14.972	1.00	31.67	T5
	ATOM	5240	O	GLU	A	82	14.655	118.226	15.378	1.00	21.74	T5
45	ATOM	5241	N	LEU	A	83	14.617	116.062	14.831	1.00	31.96	T5
	ATOM	5242	CA	LEU	A	83	16.039	115.896	15.089	1.00	25.93	T5
	ATOM	5243	CB	LEU	A	83	16.367	114.496	15.591	1.00	30.22	T5
	ATOM	5244	CG	LEU	A	83	15.887	114.141	17.000	1.00	29.55	T5
	ATOM	5245	CD1	LEU	A	83	14.364	114.144	17.047	1.00	31.12	T5
50	ATOM	5246	CD2	LEU	A	83	16.419	112.772	17.376	1.00	29.71	T5
	ATOM	5247	C	LEU	A	83	16.627	116.083	13.716	1.00	26.74	T5
	ATOM	5248	O	LEU	A	83	16.159	115.466	12.763	1.00	30.12	T5
	ATOM	5249	N	SER	A	84	17.639	116.934	13.597	1.00	28.42	T5
	ATOM	5250	CA	SER	A	84	18.227	117.201	12.291	1.00	35.60	T5
55	ATOM	5251	CB	SER	A	84	18.961	118.543	12.301	1.00	29.29	T5
	ATOM	5252	OG	SER	A	84	19.790	118.651	13.434	1.00	31.40	T5
	ATOM	5253	C	SER	A	84	19.148	116.108	11.805	1.00	27.24	T5
	ATOM	5254	O	SER	A	84	19.643	116.162	10.677	1.00	30.04	T5
	ATOM	5255	N	LEU	A	85	19.369	115.107	12.646	1.00	24.61	T5
60	ATOM	5256	CA	LEU	A	85	20.228	113.985	12.272	1.00	33.01	T5
	ATOM	5257	CB	LEU	A	85	21.393	113.866	13.268	1.00	29.56	T5
	ATOM	5258	CG	LEU	A	85	22.636	113.037	12.912	1.00	33.27	T5
	ATOM	5259	CD1	LEU	A	85	22.322	111.560	12.909	1.00	28.07	T5
	ATOM	5260	CD2	LEU	A	85	23.162	113.474	11.562	1.00	29.99	T5
65	ATOM	5261	C	LEU	A	85	19.392	112.703	12.265	1.00	30.75	T5
	ATOM	5262	O	LEU	A	85	19.002	112.194	13.315	1.00	29.74	T5

	ATOM	5263	N	VAL	A	86	19.100	112.195	11.075	1.00	30.28	T5
	ATOM	5264	CA	VAL	A	86	18.308	110.977	10.950	1.00	30.20	T5
	ATOM	5265	CB	VAL	A	86	17.171	111.121	9.914	1.00	29.02	T5
	ATOM	5266	CG1	VAL	A	86	16.462	109.793	9.749	1.00	29.59	T5
5	ATOM	5267	CG2	VAL	A	86	16.194	112.198	10.344	1.00	28.31	T5
	ATOM	5268	C	VAL	A	86	19.214	109.876	10.463	1.00	33.73	T5
	ATOM	5269	O	VAL	A	86	20.051	110.101	9.597	1.00	36.45	T5
	ATOM	5270	N	THR	A	87	19.056	108.686	11.017	1.00	29.03	T5
	ATOM	5271	CA	THR	A	87	19.878	107.587	10.572	1.00	34.21	T5
10	ATOM	5272	CB	THR	A	87	20.548	106.860	11.761	1.00	26.15	T5
	ATOM	5273	OG1	THR	A	87	20.097	105.508	11.822	1.00	33.54	T5
	ATOM	5274	CG2	THR	A	87	20.235	107.571	13.067	1.00	33.98	T5
	ATOM	5275	C	THR	A	87	19.013	106.640	9.761	1.00	28.11	T5
	ATOM	5276	O	THR	A	87	18.032	106.099	10.251	1.00	32.39	T5
15	ATOM	5277	N	LEU	A	88	19.376	106.485	8.495	1.00	32.69	T5
	ATOM	5278	CA	LEU	A	88	18.673	105.614	7.559	1.00	33.62	T5
	ATOM	5279	CB	LEU	A	88	18.714	106.221	6.157	1.00	32.27	T5
	ATOM	5280	CG	LEU	A	88	18.213	107.622	5.792	1.00	26.65	T5
	ATOM	5281	CD1	LEU	A	88	18.328	108.561	6.940	1.00	29.74	T5
20	ATOM	5282	CD2	LEU	A	88	19.022	108.143	4.641	1.00	28.45	T5
	ATOM	5283	C	LEU	A	88	19.439	104.295	7.508	1.00	27.25	T5
	ATOM	5284	O	LEU	A	88	20.670	104.269	7.644	1.00	29.28	T5
	ATOM	5285	N	PHE	A	89	18.747	103.183	7.344	1.00	34.16	T5
	ATOM	5286	CA	PHE	A	89	19.475	101.919	7.202	1.00	24.43	T5
25	ATOM	5287	CB	PHE	A	89	20.384	102.031	5.980	1.00	36.44	T5
	ATOM	5288	CG	PHE	A	89	19.715	102.702	4.825	1.00	30.39	T5
	ATOM	5289	CD1	PHE	A	89	20.433	103.518	3.964	1.00	29.83	T5
	ATOM	5290	CD2	PHE	A	89	18.323	102.588	4.651	1.00	26.14	T5
	ATOM	5291	CE1	PHE	A	89	19.772	104.228	2.946	1.00	26.70	T5
30	ATOM	5292	CE2	PHE	A	89	17.653	103.286	3.641	1.00	29.64	T5
	ATOM	5293	CZ	PHE	A	89	18.375	104.110	2.790	1.00	29.70	T5
	ATOM	5294	C	PHE	A	89	20.260	101.347	8.391	1.00	28.74	T5
	ATOM	5295	O	PHE	A	89	19.649	100.804	9.309	1.00	29.23	T5
	ATOM	5296	N	ARG	A	90	21.589	101.416	8.382	1.00	29.18	T5
35	ATOM	5297	CA	ARG	A	90	22.345	100.818	9.494	1.00	31.08	T5
	ATOM	5298	CB	ARG	A	90	21.839	101.378	10.826	1.00	36.18	T5
	ATOM	5299	CG	ARG	A	90	22.025	100.443	12.008	1.00	24.71	T5
	ATOM	5300	CD	ARG	A	90	21.168	100.863	13.175	1.00	37.20	T5
	ATOM	5301	NE	ARG	A	90	21.185	99.832	14.203	1.00	36.89	T5
40	ATOM	5302	CZ	ARG	A	90	20.139	99.507	14.957	1.00	24.97	T5
	ATOM	5303	NH1	ARG	A	90	18.974	100.141	14.807	1.00	29.81	T5
	ATOM	5304	NH2	ARG	A	90	20.250	98.522	15.845	1.00	26.25	T5
	ATOM	5305	C	ARG	A	90	22.285	99.270	9.565	1.00	30.24	T5
	ATOM	5306	O	ARG	A	90	21.207	98.672	9.571	1.00	31.80	T5
45	ATOM	5307	N	CYS	A	91	23.454	98.635	9.653	1.00	28.37	T5
	ATOM	5308	CA	CYS	A	91	23.544	97.172	9.737	1.00	24.66	T5
	ATOM	5309	CB	CYS	A	91	24.084	96.609	8.429	1.00	30.46	T5
	ATOM	5310	SG	CYS	A	91	25.655	97.335	7.988	1.00	25.00	T5
	ATOM	5311	C	CYS	A	91	24.427	96.699	10.904	1.00	32.20	T5
50	ATOM	5312	O	CYS	A	91	25.169	97.485	11.489	1.00	29.78	T5
	ATOM	5313	N	ILE	A	92	24.339	95.411	11.232	1.00	27.73	T5
	ATOM	5314	CA	ILE	A	92	25.095	94.824	12.342	1.00	31.75	T5
	ATOM	5315	CB	ILE	A	92	24.199	94.631	13.576	1.00	28.43	T5
	ATOM	5316	CG2	ILE	A	92	25.008	94.095	14.735	1.00	26.06	T5
55	ATOM	5317	CG1	ILE	A	92	23.569	95.963	13.976	1.00	28.98	T5
	ATOM	5318	CD1	ILE	A	92	22.440	95.816	14.990	1.00	27.96	T5
	ATOM	5319	C	ILE	A	92	25.629	93.453	11.969	1.00	26.62	T5
	ATOM	5320	O	ILE	A	92	25.074	92.785	11.121	1.00	29.88	T5
	ATOM	5321	N	GLN	A	93	26.710	93.030	12.606	1.00	29.74	T5
60	ATOM	5322	CA	GLN	A	93	27.287	91.717	12.331	1.00	26.65	T5
	ATOM	5323	CB	GLN	A	93	28.265	91.784	11.150	1.00	32.19	T5
	ATOM	5324	CG	GLN	A	93	27.624	91.628	9.786	1.00	28.08	T5
	ATOM	5325	CD	GLN	A	93	27.712	90.209	9.248	1.00	34.45	T5
	ATOM	5326	OE1	GLN	A	93	26.994	89.312	9.693	1.00	30.00	T5
65	ATOM	5327	NE2	GLN	A	93	28.607	89.997	8.282	1.00	37.18	T5
	ATOM	5328	C	GLN	A	93	28.014	91.177	13.558	1.00	26.29	T5

	ATOM	5329	O	GLN	A	93	28.951	91.815	14.067	1.00	25.16	T5
	ATOM	5330	N	ASN	A	94	27.574	90.017	14.044	1.00	33.86	T5
	ATOM	5331	CA	ASN	A	94	28.223	89.407	15.193	1.00	32.16	T5
	ATOM	5332	CB	ASN	A	94	27.519	88.108	15.587	1.00	29.20	T5
5	ATOM	5333	CG	ASN	A	94	26.267	88.344	16.420	1.00	34.07	T5
	ATOM	5334	OD1	ASN	A	94	26.315	89.017	17.438	1.00	34.81	T5
	ATOM	5335	ND2	ASN	A	94	25.150	87.780	15.997	1.00	26.43	T5
	ATOM	5336	C	ASN	A	94	29.656	89.118	14.764	1.00	28.81	T5
	ATOM	5337	O	ASN	A	94	29.906	88.804	13.593	1.00	17.45	T5
10	ATOM	5338	N	MET	A	95	30.598	89.245	15.693	1.00	36.76	T5
	ATOM	5339	CA	MET	A	95	31.997	88.993	15.384	1.00	37.66	T5
	ATOM	5340	CB	MET	A	95	32.853	90.207	15.759	1.00	30.87	T5
	ATOM	5341	CG	MET	A	95	32.502	91.474	15.016	1.00	37.38	T5
	ATOM	5342	SD	MET	A	95	32.453	91.239	13.219	1.00	27.32	T5
15	ATOM	5343	CE	MET	A	95	34.216	91.147	12.806	1.00	31.95	T5
	ATOM	5344	C	MET	A	95	32.508	87.775	16.146	1.00	31.49	T5
	ATOM	5345	O	MET	A	95	32.012	87.458	17.231	1.00	31.86	T5
	ATOM	5346	N	PRO	A	96	33.506	87.068	15.581	1.00	28.40	T5
	ATOM	5347	CD	PRO	A	96	34.070	87.260	14.237	1.00	29.58	T5
20	ATOM	5348	CA	PRO	A	96	34.100	85.887	16.209	1.00	28.78	T5
	ATOM	5349	CB	PRO	A	96	34.879	85.224	15.074	1.00	30.90	T5
	ATOM	5350	CG	PRO	A	96	34.330	85.839	13.828	1.00	28.73	T5
	ATOM	5351	C	PRO	A	96	35.054	86.423	17.255	1.00	32.23	T5
	ATOM	5352	O	PRO	A	96	35.080	87.626	17.532	1.00	32.88	T5
25	ATOM	5353	N	GLU	A	97	35.868	85.543	17.813	1.00	29.61	T5
	ATOM	5354	CA	GLU	A	97	36.805	85.977	18.828	1.00	37.27	T5
	ATOM	5355	CB	GLU	A	97	36.600	85.162	20.089	1.00	28.12	T5
	ATOM	5356	CG	GLU	A	97	37.241	85.782	21.289	1.00	31.00	T5
	ATOM	5357	CD	GLU	A	97	36.324	85.721	22.498	1.00	25.26	T5
30	ATOM	5358	OE1	GLU	A	97	35.956	84.589	22.907	1.00	32.01	T5
	ATOM	5359	OE2	GLU	A	97	35.961	86.801	23.037	1.00	25.27	T5
	ATOM	5360	C	GLU	A	97	38.225	85.815	18.336	1.00	31.20	T5
	ATOM	5361	O	GLU	A	97	39.145	86.486	18.814	1.00	32.99	T5
	ATOM	5362	N	THR	A	98	38.389	84.936	17.357	1.00	30.70	T5
35	ATOM	5363	CA	THR	A	98	39.698	84.653	16.811	1.00	34.40	T5
	ATOM	5364	CB	THR	A	98	39.808	83.188	16.457	1.00	20.28	T5
	ATOM	5365	OG1	THR	A	98	38.862	82.899	15.412	1.00	28.56	T5
	ATOM	5366	CG2	THR	A	98	39.501	82.326	17.685	1.00	30.44	T5
	ATOM	5367	C	THR	A	98	40.108	85.460	15.585	1.00	26.95	T5
40	ATOM	5368	O	THR	A	98	40.971	86.333	15.685	1.00	30.82	T5
	ATOM	5369	N	LEU	A	99	39.511	85.189	14.431	1.00	28.81	T5
	ATOM	5370	CA	LEU	A	99	39.935	85.914	13.243	1.00	33.27	T5
	ATOM	5371	CB	LEU	A	99	40.491	84.920	12.231	1.00	27.00	T5
	ATOM	5372	CG	LEU	A	99	41.728	84.189	12.758	1.00	27.95	T5
45	ATOM	5373	CD1	LEU	A	99	42.154	83.088	11.803	1.00	36.30	T5
	ATOM	5374	CD2	LEU	A	99	42.839	85.197	12.937	1.00	24.85	T5
	ATOM	5375	C	LEU	A	99	38.866	86.784	12.602	1.00	37.78	T5
	ATOM	5376	O	LEU	A	99	38.355	86.476	11.514	1.00	26.01	T5
	ATOM	5377	N	PRO	A	100	38.528	87.905	13.261	1.00	34.03	T5
50	ATOM	5378	CD	PRO	A	100	39.170	88.411	14.487	1.00	30.11	T5
	ATOM	5379	CA	PRO	A	100	37.514	88.850	12.782	1.00	31.85	T5
	ATOM	5380	CB	PRO	A	100	37.748	90.067	13.662	1.00	20.86	T5
	ATOM	5381	CG	PRO	A	100	38.170	89.441	14.961	1.00	28.49	T5
	ATOM	5382	C	PRO	A	100	37.637	89.175	11.300	1.00	35.91	T5
55	ATOM	5383	O	PRO	A	100	38.674	89.642	10.834	1.00	33.87	T5
	ATOM	5384	N	ASN	A	101	36.563	88.921	10.571	1.00	29.55	T5
	ATOM	5385	CA	ASN	A	101	36.517	89.181	9.145	1.00	38.56	T5
	ATOM	5386	CB	ASN	A	101	37.282	88.117	8.389	1.00	30.68	T5
	ATOM	5387	CG	ASN	A	101	38.721	88.445	8.258	1.00	25.57	T5
60	ATOM	5388	OD1	ASN	A	101	39.089	89.379	7.540	1.00	35.15	T5
	ATOM	5389	ND2	ASN	A	101	39.569	87.686	8.957	1.00	28.07	T5
	ATOM	5390	C	ASN	A	101	35.091	89.142	8.671	1.00	24.27	T5
	ATOM	5391	O	ASN	A	101	34.645	88.113	8.162	1.00	29.66	T5
	ATOM	5392	N	ASN	A	102	34.358	90.240	8.811	1.00	23.87	T5
65	ATOM	5393	CA	ASN	A	102	32.991	90.166	8.361	1.00	31.88	T5
	ATOM	5394	CB	ASN	A	102	32.040	90.232	9.552	1.00	20.87	T5

	ATOM	5395	CG	ASN	A	102	31.851	88.860	10.196	1.00	27.88	T5
	ATOM	5396	OD1	ASN	A	102	31.666	87.851	9.498	1.00	34.35	T5
	ATOM	5397	ND2	ASN	A	102	31.904	88.812	11.522	1.00	19.93	T5
	ATOM	5398	C	ASN	A	102	32.487	91.012	7.212	1.00	33.15	T5
5	ATOM	5399	O	ASN	A	102	31.848	90.459	6.323	1.00	31.78	T5
	ATOM	5400	N	SER	A	103	32.744	92.308	7.158	1.00	34.77	T5
	ATOM	5401	CA	SER	A	103	32.209	93.037	6.001	1.00	27.16	T5
	ATOM	5402	CB	SER	A	103	32.789	92.465	4.701	1.00	21.65	T5
	ATOM	5403	OG	SER	A	103	31.772	92.228	3.740	1.00	26.80	T5
10	ATOM	5404	C	SER	A	103	30.672	92.961	5.944	1.00	28.56	T5
	ATOM	5405	O	SER	A	103	30.064	91.888	6.016	1.00	30.58	T5
	ATOM	5406	N	CYS	A	104	30.034	94.109	5.800	1.00	30.93	T5
	ATOM	5407	CA	CYS	A	104	28.589	94.126	5.764	1.00	33.70	T5
	ATOM	5408	CB	CYS	A	104	28.057	94.337	7.179	1.00	30.83	T5
15	ATOM	5409	SG	CYS	A	104	26.294	94.063	7.358	1.00	33.35	T5
	ATOM	5410	C	CYS	A	104	28.113	95.230	4.840	1.00	37.34	T5
	ATOM	5411	O	CYS	A	104	28.532	96.378	4.963	1.00	30.02	T5
	ATOM	5412	N	TYR	A	105	27.248	94.864	3.905	1.00	41.35	T5
	ATOM	5413	CA	TYR	A	105	26.697	95.805	2.947	1.00	31.33	T5
20	ATOM	5414	CB	TYR	A	105	26.777	95.215	1.540	1.00	33.17	T5
	ATOM	5415	CG	TYR	A	105	26.139	96.061	0.457	1.00	30.86	T5
	ATOM	5416	CD1	TYR	A	105	26.895	96.950	-0.296	1.00	29.25	T5
	ATOM	5417	CE1	TYR	A	105	26.315	97.711	-1.310	1.00	27.99	T5
	ATOM	5418	CD2	TYR	A	105	24.779	95.959	0.173	1.00	29.43	T5
25	ATOM	5419	CE2	TYR	A	105	24.191	96.722	-0.836	1.00	29.77	T5
	ATOM	5420	CZ	TYR	A	105	24.969	97.589	-1.570	1.00	33.16	T5
	ATOM	5421	OH	TYR	A	105	24.412	98.321	-2.581	1.00	24.48	T5
	ATOM	5422	C	TYR	A	105	25.243	96.084	3.286	1.00	33.42	T5
	ATOM	5423	O	TYR	A	105	24.532	95.211	3.776	1.00	33.88	T5
30	ATOM	5424	N	SER	A	106	24.804	97.306	3.029	1.00	34.28	T5
	ATOM	5425	CA	SER	A	106	23.422	97.667	3.279	1.00	30.49	T5
	ATOM	5426	CB	SER	A	106	23.203	97.974	4.760	1.00	30.07	T5
	ATOM	5427	OG	SER	A	106	21.825	98.081	5.062	1.00	27.17	T5
	ATOM	5428	C	SER	A	106	23.138	98.887	2.424	1.00	31.93	T5
35	ATOM	5429	O	SER	A	106	24.021	99.717	2.226	1.00	29.76	T5
	ATOM	5430	N	ALA	A	107	21.917	98.980	1.901	1.00	31.42	T5
	ATOM	5431	CA	ALA	A	107	21.527	100.100	1.054	1.00	31.87	T5
	ATOM	5432	CB	ALA	A	107	21.995	99.853	-0.375	1.00	25.74	T5
	ATOM	5433	C	ALA	A	107	20.021	100.313	1.079	1.00	29.56	T5
40	ATOM	5434	O	ALA	A	107	19.271	99.449	1.511	1.00	29.94	T5
	ATOM	5435	N	GLY	A	108	19.589	101.473	0.605	1.00	28.67	T5
	ATOM	5436	CA	GLY	A	108	18.174	101.783	0.570	1.00	38.52	T5
	ATOM	5437	C	GLY	A	108	17.910	103.070	-0.184	1.00	36.57	T5
	ATOM	5438	O	GLY	A	108	18.840	103.706	-0.667	1.00	31.53	T5
45	ATOM	5439	N	ILE	A	109	16.642	103.453	-0.291	1.00	31.45	T5
	ATOM	5440	CA	ILE	A	109	16.264	104.678	-0.987	1.00	29.07	T5
	ATOM	5441	CB	ILE	A	109	15.168	104.413	-2.029	1.00	31.26	T5
	ATOM	5442	CG2	ILE	A	109	14.784	105.700	-2.712	1.00	27.53	T5
	ATOM	5443	CG1	ILE	A	109	15.664	103.409	-3.058	1.00	33.79	T5
50	ATOM	5444	CD1	ILE	A	109	14.603	102.933	-3.992	1.00	23.60	T5
	ATOM	5445	C	ILE	A	109	15.731	105.687	0.023	1.00	30.13	T5
	ATOM	5446	O	ILE	A	109	15.038	105.318	0.972	1.00	36.96	T5
	ATOM	5447	N	ALA	A	110	16.064	106.958	-0.173	1.00	27.02	T5
	ATOM	5448	CA	ALA	A	110	15.601	108.004	0.725	1.00	28.41	T5
55	ATOM	5449	CB	ALA	A	110	16.585	108.200	1.840	1.00	32.27	T5
	ATOM	5450	C	ALA	A	110	15.443	109.291	-0.056	1.00	36.92	T5
	ATOM	5451	O	ALA	A	110	16.068	109.469	-1.102	1.00	31.91	T5
	ATOM	5452	N	LYS	A	111	14.588	110.184	0.433	1.00	23.81	T5
	ATOM	5453	CA	LYS	A	111	14.399	111.455	-0.246	1.00	26.15	T5
60	ATOM	5454	CB	LYS	A	111	12.952	111.914	-0.175	1.00	31.15	T5
	ATOM	5455	CG	LYS	A	111	12.738	113.215	-0.934	1.00	29.11	T5
	ATOM	5456	CD	LYS	A	111	11.280	113.660	-0.917	1.00	27.86	T5
	ATOM	5457	CE	LYS	A	111	11.083	114.955	-1.717	1.00	34.10	T5
	ATOM	5458	NZ	LYS	A	111	9.638	115.394	-1.737	1.00	25.58	T5
65	ATOM	5459	C	LYS	A	111	15.286	112.461	0.453	1.00	28.57	T5
	ATOM	5460	O	LYS	A	111	15.276	112.541	1.681	1.00	34.31	T5

	ATOM	5461	N	LEU	A	112	16.052	113.218	-0.330	1.00	30.53	T5
	ATOM	5462	CA	LEU	A	112	16.986	114.206	0.209	1.00	31.60	T5
	ATOM	5463	CB	LEU	A	112	18.426	113.755	-0.060	1.00	35.58	T5
	ATOM	5464	CG	LEU	A	112	18.791	112.319	0.315	1.00	28.84	T5
5	ATOM	5465	CD1	LEU	A	112	20.119	111.955	-0.291	1.00	33.34	T5
	ATOM	5466	CD2	LEU	A	112	18.814	112.177	1.823	1.00	33.23	T5
	ATOM	5467	C	LEU	A	112	16.770	115.570	-0.436	1.00	33.98	T5
	ATOM	5468	O	LEU	A	112	16.170	115.658	-1.501	1.00	33.90	T5
	ATOM	5469	N	GLU	A	113	17.273	116.623	0.206	1.00	28.73	T5
10	ATOM	5470	CA	GLU	A	113	17.144	117.976	-0.318	1.00	37.12	T5
	ATOM	5471	CB	GLU	A	113	16.250	118.826	0.564	1.00	26.04	T5
	ATOM	5472	CG	GLU	A	113	15.053	118.163	1.135	1.00	25.55	T5
	ATOM	5473	CD	GLU	A	113	14.018	119.187	1.586	1.00	37.32	T5
	ATOM	5474	OE1	GLU	A	113	14.406	120.195	2.247	1.00	27.52	T5
15	ATOM	5475	OE2	GLU	A	113	12.811	118.977	1.272	1.00	30.85	T5
	ATOM	5476	C	GLU	A	113	18.474	118.705	-0.395	1.00	36.46	T5
	ATOM	5477	O	GLU	A	113	19.424	118.366	0.315	1.00	24.79	T5
	ATOM	5478	N	GLU	A	114	18.520	119.731	-1.243	1.00	28.25	T5
	ATOM	5479	CA	GLU	A	114	19.710	120.559	-1.391	1.00	28.32	T5
20	ATOM	5480	CB	GLU	A	114	19.371	121.882	-2.039	1.00	34.65	T5
	ATOM	5481	CG	GLU	A	114	19.431	121.924	-3.522	1.00	33.64	T5
	ATOM	5482	CD	GLU	A	114	19.535	123.357	-3.997	1.00	33.43	T5
	ATOM	5483	OE1	GLU	A	114	20.565	124.008	-3.688	1.00	37.93	T5
	ATOM	5484	OE2	GLU	A	114	18.586	123.834	-4.660	1.00	30.69	T5
25	ATOM	5485	C	GLU	A	114	20.263	120.901	-0.028	1.00	35.46	T5
	ATOM	5486	O	GLU	A	114	19.549	121.451	0.803	1.00	32.15	T5
	ATOM	5487	N	GLY	A	115	21.532	120.602	0.200	1.00	30.65	T5
	ATOM	5488	CA	GLY	A	115	22.123	120.937	1.478	1.00	36.03	T5
	ATOM	5489	C	GLY	A	115	22.208	119.791	2.456	1.00	31.08	T5
30	ATOM	5490	O	GLY	A	115	22.958	119.875	3.428	1.00	32.33	T5
	ATOM	5491	N	ASP	A	116	21.436	118.732	2.235	1.00	28.10	T5
	ATOM	5492	CA	ASP	A	116	21.513	117.599	3.142	1.00	31.21	T5
	ATOM	5493	CB	ASP	A	116	20.456	116.541	2.807	1.00	28.60	T5
	ATOM	5494	CG	ASP	A	116	19.050	116.973	3.165	1.00	33.01	T5
35	ATOM	5495	OD1	ASP	A	116	18.896	117.819	4.067	1.00	31.09	T5
	ATOM	5496	OD2	ASP	A	116	18.097	116.445	2.556	1.00	28.94	T5
	ATOM	5497	C	ASP	A	116	22.901	116.983	2.979	1.00	28.02	T5
	ATOM	5498	O	ASP	A	116	23.524	117.108	1.918	1.00	33.63	T5
	ATOM	5499	N	GLU	A	117	23.393	116.334	4.028	1.00	33.45	T5
40	ATOM	5500	CA	GLU	A	117	24.690	115.691	3.947	1.00	32.25	T5
	ATOM	5501	CB	GLU	A	117	25.695	116.391	4.847	1.00	36.41	T5
	ATOM	5502	CG	GLU	A	117	25.828	117.870	4.586	1.00	37.23	T5
	ATOM	5503	CD	GLU	A	117	26.915	118.508	5.433	1.00	24.48	T5
	ATOM	5504	OE1	GLU	A	117	27.064	118.109	6.615	1.00	26.88	T5
45	ATOM	5505	OE2	GLU	A	117	27.611	119.416	4.926	1.00	31.95	T5
	ATOM	5506	C	GLU	A	117	24.533	114.254	4.388	1.00	29.17	T5
	ATOM	5507	O	GLU	A	117	23.772	113.975	5.316	1.00	30.06	T5
	ATOM	5508	N	LEU	A	118	25.234	113.349	3.707	1.00	35.25	T5
	ATOM	5509	CA	LEU	A	118	25.201	111.932	4.042	1.00	30.55	T5
50	ATOM	5510	CB	LEU	A	118	24.957	111.084	2.796	1.00	29.96	T5
	ATOM	5511	CG	LEU	A	118	23.605	111.236	2.109	1.00	36.10	T5
	ATOM	5512	CD1	LEU	A	118	23.514	110.267	0.966	1.00	29.52	T5
	ATOM	5513	CD2	LEU	A	118	22.491	110.973	3.104	1.00	31.15	T5
	ATOM	5514	C	LEU	A	118	26.538	111.552	4.643	1.00	29.68	T5
55	ATOM	5515	O	LEU	A	118	27.565	112.085	4.246	1.00	31.32	T5
	ATOM	5516	N	GLN	A	119	26.526	110.639	5.605	1.00	32.14	T5
	ATOM	5517	CA	GLN	A	119	27.758	110.190	6.224	1.00	34.31	T5
	ATOM	5518	CB	GLN	A	119	28.120	111.108	7.375	1.00	31.10	T5
	ATOM	5519	CG	GLN	A	119	27.124	111.107	8.495	1.00	31.65	T5
60	ATOM	5520	CD	GLN	A	119	27.480	112.099	9.580	1.00	31.79	T5
	ATOM	5521	OE1	GLN	A	119	26.980	112.012	10.700	1.00	29.92	T5
	ATOM	5522	NE2	GLN	A	119	28.338	113.057	9.250	1.00	26.45	T5
	ATOM	5523	C	GLN	A	119	27.639	108.749	6.714	1.00	31.78	T5
	ATOM	5524	O	GLN	A	119	26.545	108.268	6.999	1.00	28.96	T5
65	ATOM	5525	N	LEU	A	120	28.778	108.068	6.797	1.00	31.82	T5
	ATOM	5526	CA	LEU	A	120	28.857	106.673	7.240	1.00	21.13	T5

	ATOM	5527	CB	LEU	A	120	29.710	105.887	6.241	1.00	27.50	T5
	ATOM	5528	CG	LEU	A	120	29.822	104.361	6.195	1.00	28.20	T5
	ATOM	5529	CD1	LEU	A	120	30.027	103.796	7.586	1.00	34.87	T5
	ATOM	5530	CD2	LEU	A	120	28.579	103.799	5.551	1.00	31.02	T5
5	ATOM	5531	C	LEU	A	120	29.534	106.672	8.614	1.00	27.45	T5
	ATOM	5532	O	LEU	A	120	30.670	107.134	8.745	1.00	29.28	T5
	ATOM	5533	N	ALA	A	121	28.856	106.155	9.635	1.00	29.57	T5
	ATOM	5534	CA	ALA	A	121	29.434	106.156	10.977	1.00	26.75	T5
	ATOM	5535	CB	ALA	A	121	28.723	107.181	11.836	1.00	34.19	T5
10	ATOM	5536	C	ALA	A	121	29.446	104.816	11.697	1.00	33.27	T5
	ATOM	5537	O	ALA	A	121	28.503	104.038	11.608	1.00	34.23	T5
	ATOM	5538	N	ILE	A	122	30.537	104.558	12.412	1.00	32.62	T5
	ATOM	5539	CA	ILE	A	122	30.686	103.331	13.187	1.00	30.40	T5
	ATOM	5540	CB	ILE	A	122	32.044	102.666	12.943	1.00	36.82	T5
15	ATOM	5541	CG2	ILE	A	122	32.121	101.378	13.719	1.00	40.91	T5
	ATOM	5542	CG1	ILE	A	122	32.215	102.369	11.460	1.00	26.12	T5
	ATOM	5543	CD1	ILE	A	122	33.514	101.714	11.109	1.00	29.29	T5
	ATOM	5544	C	ILE	A	122	30.569	103.693	14.667	1.00	26.96	T5
	ATOM	5545	O	ILE	A	122	31.375	104.462	15.204	1.00	31.08	T5
20	ATOM	5546	N	PRO	A	123	29.549	103.146	15.343	1.00	33.89	T5
	ATOM	5547	CD	PRO	A	123	28.487	102.323	14.750	1.00	36.39	T5
	ATOM	5548	CA	PRO	A	123	29.281	103.391	16.765	1.00	34.34	T5
	ATOM	5549	CB	PRO	A	123	27.872	102.834	16.962	1.00	31.76	T5
	ATOM	5550	CG	PRO	A	123	27.312	102.749	15.567	1.00	25.80	T5
25	ATOM	5551	C	PRO	A	123	30.283	102.725	17.710	1.00	27.83	T5
	ATOM	5552	O	PRO	A	123	29.892	101.968	18.599	1.00	26.29	T5
	ATOM	5553	N	ARG	A	124	31.570	102.994	17.511	1.00	28.02	T5
	ATOM	5554	CA	ARG	A	124	32.610	102.437	18.365	1.00	23.25	T5
	ATOM	5555	CB	ARG	A	124	33.143	101.135	17.810	1.00	35.72	T5
30	ATOM	5556	CG	ARG	A	124	32.339	99.951	18.233	1.00	40.64	T5
	ATOM	5557	CD	ARG	A	124	33.275	98.836	18.623	1.00	28.23	T5
	ATOM	5558	NE	ARG	A	124	33.935	99.090	19.908	1.00	28.54	T5
	ATOM	5559	CZ	ARG	A	124	34.937	98.349	20.385	1.00	29.58	T5
	ATOM	5560	NH1	ARG	A	124	35.392	97.318	19.680	1.00	24.30	T5
35	ATOM	5561	NH2	ARG	A	124	35.475	98.622	21.569	1.00	33.53	T5
	ATOM	5562	C	ARG	A	124	33.736	103.427	18.460	1.00	28.72	T5
	ATOM	5563	O	ARG	A	124	33.903	104.258	17.565	1.00	26.37	T5
	ATOM	5564	N	GLU	A	125	34.515	103.339	19.536	1.00	26.24	T5
	ATOM	5565	CA	GLU	A	125	35.607	104.273	19.726	1.00	34.88	T5
40	ATOM	5566	CB	GLU	A	125	36.069	104.251	21.168	1.00	34.15	T5
	ATOM	5567	CG	GLU	A	125	35.067	104.910	22.092	1.00	31.19	T5
	ATOM	5568	CD	GLU	A	125	35.743	105.789	23.148	1.00	29.71	T5
	ATOM	5569	OE1	GLU	A	125	36.613	105.267	23.907	1.00	28.89	T5
	ATOM	5570	OE2	GLU	A	125	35.403	107.001	23.218	1.00	26.43	T5
45	ATOM	5571	C	GLU	A	125	36.783	104.098	18.779	1.00	26.44	T5
	ATOM	5572	O	GLU	A	125	37.219	105.073	18.163	1.00	31.41	T5
	ATOM	5573	N	ASN	A	126	37.315	102.890	18.653	1.00	26.87	T5
	ATOM	5574	CA	ASN	A	126	38.414	102.705	17.717	1.00	33.12	T5
	ATOM	5575	CB	ASN	A	126	39.758	102.663	18.432	1.00	25.75	T5
50	ATOM	5576	CG	ASN	A	126	40.288	104.052	18.765	1.00	31.34	T5
	ATOM	5577	OD1	ASN	A	126	39.867	104.676	19.744	1.00	22.21	T5
	ATOM	5578	ND2	ASN	A	126	41.215	104.546	17.939	1.00	27.25	T5
	ATOM	5579	C	ASN	A	126	38.185	101.429	16.960	1.00	25.41	T5
	ATOM	5580	O	ASN	A	126	38.953	100.474	17.058	1.00	32.69	T5
55	ATOM	5581	N	ALA	A	127	37.096	101.427	16.207	1.00	35.95	T5
	ATOM	5582	CA	ALA	A	127	36.707	100.278	15.421	1.00	28.70	T5
	ATOM	5583	CB	ALA	A	127	35.580	100.668	14.481	1.00	29.29	T5
	ATOM	5584	C	ALA	A	127	37.871	99.709	14.631	1.00	33.39	T5
	ATOM	5585	O	ALA	A	127	38.613	100.453	13.993	1.00	32.28	T5
60	ATOM	5586	N	GLN	A	128	38.039	98.390	14.701	1.00	24.24	T5
	ATOM	5587	CA	GLN	A	128	39.091	97.710	13.949	1.00	34.44	T5
	ATOM	5588	CB	GLN	A	128	39.474	96.402	14.638	1.00	24.62	T5
	ATOM	5589	CG	GLN	A	128	40.142	96.621	15.992	1.00	29.66	T5
	ATOM	5590	CD	GLN	A	128	41.217	97.698	15.919	1.00	27.41	T5
65	ATOM	5591	OE1	GLN	A	128	42.167	97.594	15.140	1.00	27.50	T5
	ATOM	5592	NE2	GLN	A	128	41.062	98.745	16.724	1.00	32.06	T5

	ATOM	5593	C	GLN	A	128	38.493	97.453	12.574	1.00	28.52	T5
	ATOM	5594	O	GLN	A	128	37.721	96.513	12.373	1.00	30.38	T5
	ATOM	5595	N	ILE	A	129	38.866	98.307	11.631	1.00	41.09	T5
	ATOM	5596	CA	ILE	A	129	38.333	98.282	10.280	1.00	27.24	T5
5	ATOM	5597	CB	ILE	A	129	37.555	99.628	10.062	1.00	27.72	T5
	ATOM	5598	CG2	ILE	A	129	37.977	100.335	8.796	1.00	34.86	T5
	ATOM	5599	CG1	ILE	A	129	36.061	99.357	10.116	1.00	24.64	T5
	ATOM	5600	CD1	ILE	A	129	35.617	98.776	11.441	1.00	29.52	T5
	ATOM	5601	C	ILE	A	129	39.388	98.086	9.204	1.00	29.37	T5
10	ATOM	5602	O	ILE	A	129	40.565	98.328	9.437	1.00	30.77	T5
	ATOM	5603	N	SER	A	130	38.972	97.626	8.031	1.00	27.57	T5
	ATOM	5604	CA	SER	A	130	39.901	97.469	6.922	1.00	29.52	T5
	ATOM	5605	CB	SER	A	130	39.622	96.187	6.154	1.00	33.63	T5
	ATOM	5606	OG	SER	A	130	40.340	96.188	4.935	1.00	30.42	T5
15	ATOM	5607	C	SER	A	130	39.686	98.658	5.997	1.00	27.44	T5
	ATOM	5608	O	SER	A	130	38.564	98.914	5.583	1.00	28.70	T5
	ATOM	5609	N	LEU	A	131	40.743	99.388	5.670	1.00	28.04	T5
	ATOM	5610	CA	LEU	A	131	40.590	100.541	4.791	1.00	29.67	T5
	ATOM	5611	CB	LEU	A	131	41.434	101.709	5.295	1.00	30.23	T5
20	ATOM	5612	CG	LEU	A	131	40.846	102.494	6.463	1.00	39.51	T5
	ATOM	5613	CD1	LEU	A	131	40.623	101.591	7.636	1.00	30.02	T5
	ATOM	5614	CD2	LEU	A	131	41.794	103.607	6.837	1.00	29.03	T5
	ATOM	5615	C	LEU	A	131	40.919	100.275	3.328	1.00	32.44	T5
	ATOM	5616	O	LEU	A	131	41.412	101.158	2.629	1.00	32.59	T5
25	ATOM	5617	N	ASP	A	132	40.638	99.065	2.859	1.00	30.60	T5
	ATOM	5618	CA	ASP	A	132	40.896	98.727	1.464	1.00	30.10	T5
	ATOM	5619	CB	ASP	A	132	41.166	97.228	1.320	1.00	32.15	T5
	ATOM	5620	CG	ASP	A	132	42.599	96.862	1.641	1.00	31.33	T5
	ATOM	5621	OD1	ASP	A	132	42.882	95.652	1.790	1.00	35.82	T5
30	ATOM	5622	OD2	ASP	A	132	43.440	97.783	1.734	1.00	29.49	T5
	ATOM	5623	C	ASP	A	132	39.748	99.136	0.541	1.00	32.84	T5
	ATOM	5624	O	ASP	A	132	38.569	99.019	0.889	1.00	34.45	T5
	ATOM	5625	N	GLY	A	133	40.110	99.617	-0.642	1.00	36.40	T5
	ATOM	5626	CA	GLY	A	133	39.121	100.041	-1.612	1.00	28.36	T5
35	ATOM	5627	C	GLY	A	133	38.045	99.023	-1.947	1.00	26.24	T5
	ATOM	5628	O	GLY	A	133	36.919	99.418	-2.237	1.00	29.63	T5
	ATOM	5629	N	ASP	A	134	38.369	97.728	-1.916	1.00	26.16	T5
	ATOM	5630	CA	ASP	A	134	37.381	96.700	-2.234	1.00	29.34	T5
	ATOM	5631	CB	ASP	A	134	37.977	95.305	-2.312	1.00	35.22	T5
40	ATOM	5632	CG	ASP	A	134	39.404	95.303	-2.648	1.00	27.26	T5
	ATOM	5633	OD1	ASP	A	134	39.719	95.762	-3.754	1.00	32.75	T5
	ATOM	5634	OD2	ASP	A	134	40.203	94.834	-1.812	1.00	30.75	T5
	ATOM	5635	C	ASP	A	134	36.348	96.577	-1.155	1.00	30.33	T5
	ATOM	5636	O	ASP	A	134	35.160	96.697	-1.391	1.00	32.28	T5
45	ATOM	5637	N	VAL	A	135	36.832	96.283	0.037	1.00	29.19	T5
	ATOM	5638	CA	VAL	A	135	35.977	96.044	1.172	1.00	33.63	T5
	ATOM	5639	CB	VAL	A	135	36.790	95.384	2.263	1.00	26.73	T5
	ATOM	5640	CG1	VAL	A	135	37.340	94.082	1.737	1.00	28.77	T5
	ATOM	5641	CG2	VAL	A	135	37.922	96.295	2.677	1.00	26.40	T5
50	ATOM	5642	C	VAL	A	135	35.132	97.158	1.772	1.00	26.11	T5
	ATOM	5643	O	VAL	A	135	34.009	96.891	2.190	1.00	30.47	T5
	ATOM	5644	N	THR	A	136	35.626	98.390	1.843	1.00	27.05	T5
	ATOM	5645	CA	THR	A	136	34.783	99.431	2.424	1.00	30.39	T5
	ATOM	5646	CB	THR	A	136	35.295	99.859	3.826	1.00	27.49	T5
55	ATOM	5647	OG1	THR	A	136	36.427	100.712	3.686	1.00	32.44	T5
	ATOM	5648	CG2	THR	A	136	35.711	98.643	4.635	1.00	39.73	T5
	ATOM	5649	C	THR	A	136	34.606	100.652	1.515	1.00	30.10	T5
	ATOM	5650	O	THR	A	136	35.573	101.313	1.128	1.00	34.47	T5
	ATOM	5651	N	PHE	A	137	33.349	100.934	1.170	1.00	30.68	T5
60	ATOM	5652	CA	PHE	A	137	33.010	102.048	0.293	1.00	32.14	T5
	ATOM	5653	CB	PHE	A	137	32.998	101.550	-1.145	1.00	27.33	T5
	ATOM	5654	CG	PHE	A	137	32.303	100.234	-1.323	1.00	29.61	T5
	ATOM	5655	CD1	PHE	A	137	30.922	100.161	-1.414	1.00	33.19	T5
	ATOM	5656	CD2	PHE	A	137	33.031	99.067	-1.411	1.00	26.04	T5
65	ATOM	5657	CE1	PHE	A	137	30.283	98.941	-1.593	1.00	33.37	T5
	ATOM	5658	CE2	PHE	A	137	32.397	97.843	-1.588	1.00	30.02	T5

	ATOM	5659	CZ	PHE A 137	31.026	97.783	-1.680	1.00	31.61	T5
	ATOM	5660	C	PHE A 137	31.670	102.702	0.658	1.00	30.40	T5
	ATOM	5661	O	PHE A 137	30.871	102.122	1.395	1.00	35.93	T5
	ATOM	5662	N	PHE A 138	31.427	103.903	0.133	1.00	31.92	T5
5	ATOM	5663	CA	PHE A 138	30.207	104.642	0.444	1.00	24.12	T5
	ATOM	5664	CB	PHE A 138	30.591	105.823	1.333	1.00	35.08	T5
	ATOM	5665	CG	PHE A 138	29.438	106.490	2.009	1.00	34.39	T5
	ATOM	5666	CD1	PHE A 138	28.245	105.818	2.218	1.00	25.97	T5
	ATOM	5667	CD2	PHE A 138	29.551	107.803	2.444	1.00	25.30	T5
10	ATOM	5668	CE1	PHE A 138	27.177	106.449	2.848	1.00	29.80	T5
	ATOM	5669	CE2	PHE A 138	28.492	108.437	3.074	1.00	34.01	T5
	ATOM	5670	CZ	PHE A 138	27.303	107.759	3.275	1.00	28.28	T5
	ATOM	5671	C	PHE A 138	29.445	105.080	-0.817	1.00	28.29	T5
	ATOM	5672	O	PHE A 138	30.013	105.655	-1.742	1.00	31.28	T5
15	ATOM	5673	N	GLY A 139	28.141	104.804	-0.806	1.00	34.83	T5
	ATOM	5674	CA	GLY A 139	27.237	105.044	-1.933	1.00	32.28	T5
	ATOM	5675	C	GLY A 139	26.712	106.366	-2.452	1.00	37.53	T5
	ATOM	5676	O	GLY A 139	27.477	107.288	-2.649	1.00	31.17	T5
	ATOM	5677	N	ALA A 140	25.407	106.416	-2.725	1.00	31.72	T5
20	ATOM	5678	CA	ALA A 140	24.693	107.604	-3.251	1.00	35.24	T5
	ATOM	5679	CB	ALA A 140	25.265	108.870	-2.660	1.00	27.65	T5
	ATOM	5680	C	ALA A 140	24.591	107.763	-4.782	1.00	29.75	T5
	ATOM	5681	O	ALA A 140	25.539	108.143	-5.451	1.00	26.48	T5
	ATOM	5682	N	LEU A 141	23.405	107.485	-5.310	1.00	28.78	T5
25	ATOM	5683	CA	LEU A 141	23.102	107.583	-6.735	1.00	26.18	T5
	ATOM	5684	CB	LEU A 141	23.146	106.195	-7.379	1.00	33.59	T5
	ATOM	5685	CG	LEU A 141	22.684	106.043	-8.831	1.00	32.90	T5
	ATOM	5686	CD1	LEU A 141	23.185	104.748	-9.407	1.00	29.28	T5
	ATOM	5687	CD2	LEU A 141	21.181	106.062	-8.895	1.00	27.07	T5
30	ATOM	5688	C	LEU A 141	21.697	108.175	-6.874	1.00	29.23	T5
	ATOM	5689	O	LEU A 141	20.766	107.747	-6.199	1.00	27.27	T5
	ATOM	5690	N	LYS A 142	21.533	109.148	-7.761	1.00	26.56	T5
	ATOM	5691	CA	LYS A 142	20.228	109.776	-7.937	1.00	26.15	T5
	ATOM	5692	CB	LYS A 142	20.406	111.234	-8.362	1.00	25.63	T5
35	ATOM	5693	CG	LYS A 142	19.084	111.948	-8.550	1.00	26.74	T5
	ATOM	5694	CD	LYS A 142	19.238	113.439	-8.680	1.00	24.13	T5
	ATOM	5695	CE	LYS A 142	17.871	114.078	-8.841	1.00	33.96	T5
	ATOM	5696	NZ	LYS A 142	17.962	115.560	-9.001	1.00	30.38	T5
	ATOM	5697	C	LYS A 142	19.275	109.080	-8.914	1.00	32.61	T5
40	ATOM	5698	O	LYS A 142	19.631	108.790	-10.056	1.00	27.71	T5
	ATOM	5699	N	LEU A 143	18.052	108.829	-8.457	1.00	31.32	T5
	ATOM	5700	CA	LEU A 143	17.040	108.184	-9.287	1.00	30.86	T5
	ATOM	5701	CB	LEU A 143	15.974	107.535	-8.403	1.00	29.46	T5
	ATOM	5702	CG	LEU A 143	16.412	106.480	-7.386	1.00	28.20	T5
45	ATOM	5703	CD1	LEU A 143	15.211	106.045	-6.576	1.00	26.77	T5
	ATOM	5704	CD2	LEU A 143	17.031	105.290	-8.088	1.00	33.54	T5
	ATOM	5705	C	LEU A 143	16.369	109.205	-10.207	1.00	29.78	T5
	ATOM	5706	O	LEU A 143	16.339	110.399	-9.903	1.00	30.07	T5
	ATOM	5707	N	LEU A 144	15.828	108.741	-11.330	1.00	23.66	T5
50	ATOM	5708	CA	LEU A 144	15.143	109.640	-12.259	1.00	24.84	T5
	ATOM	5709	CB	LEU A 144	15.218	109.100	-13.683	1.00	28.37	T5
	ATOM	5710	CG	LEU A 144	16.613	109.049	-14.293	1.00	34.85	T5
	ATOM	5711	CD1	LEU A 144	16.543	108.322	-15.608	1.00	30.05	T5
	ATOM	5712	CD2	LEU A 144	17.162	110.456	-14.479	1.00	30.03	T5
55	ATOM	5713	C	LEU A 144	13.677	109.784	-11.863	1.00	25.99	T5
	ATOM	5714	O	LEU A 144	13.171	108.927	-11.105	1.00	28.47	T5
	ATOM	5715	OXT	LEU A 144	13.041	110.748	-12.333	1.00	24.27	T5
	ATOM	5716	CB	VAL A 1	9.357	110.997	-17.279	1.00	33.27	T6
	ATOM	5717	CG1	VAL A 1	9.972	112.323	-16.795	1.00	32.40	T6
60	ATOM	5718	CG2	VAL A 1	8.662	111.204	-18.631	1.00	35.51	T6
	ATOM	5719	C	VAL A 1	11.049	109.617	-16.003	1.00	30.71	T6
	ATOM	5720	O	VAL A 1	12.242	109.836	-15.789	1.00	34.57	T6
	ATOM	5721	N	VAL A 1	9.942	108.638	-18.032	1.00	24.53	T6
	ATOM	5722	CA	VAL A 1	10.470	109.887	-17.401	1.00	24.55	T6
65	ATOM	5723	N	THR A 2	10.225	109.148	-15.059	1.00	33.67	T6
	ATOM	5724	CA	THR A 2	10.716	108.879	-13.704	1.00	30.57	T6

	ATOM	5725	CB	THR	A	2	9.886	109.607	-12.642	1.00	32.32	T6
	ATOM	5726	OG1	THR	A	2	8.597	108.986	-12.542	1.00	29.88	T6
	ATOM	5727	CG2	THR	A	2	9.734	111.082	-13.002	1.00	41.52	T6
	ATOM	5728	C	THR	A	2	10.707	107.401	-13.344	1.00	31.56	T6
5	ATOM	5729	O	THR	A	2	10.176	106.582	-14.092	1.00	26.14	T6
	ATOM	5730	N	GLN	A	3	11.284	107.070	-12.188	1.00	32.06	T6
	ATOM	5731	CA	GLN	A	3	11.347	105.685	-11.725	1.00	30.58	T6
	ATOM	5732	CB	GLN	A	3	12.789	105.301	-11.405	1.00	27.93	T6
	ATOM	5733	CG	GLN	A	3	13.789	105.705	-12.459	1.00	34.17	T6
10	ATOM	5734	CD	GLN	A	3	15.170	105.161	-12.170	1.00	26.23	T6
	ATOM	5735	OE1	GLN	A	3	15.358	103.948	-12.104	1.00	33.74	T6
	ATOM	5736	NE2	GLN	A	3	16.145	106.053	-11.994	1.00	34.63	T6
	ATOM	5737	C	GLN	A	3	10.500	105.461	-10.482	1.00	33.30	T6
	ATOM	5738	O	GLN	A	3	10.887	105.858	-9.382	1.00	29.18	T6
15	ATOM	5739	N	ASP	A	4	9.351	104.818	-10.645	1.00	29.79	T6
	ATOM	5740	CA	ASP	A	4	8.500	104.562	-9.495	1.00	27.72	T6
	ATOM	5741	CB	ASP	A	4	7.198	103.886	-9.923	1.00	31.38	T6
	ATOM	5742	CG	ASP	A	4	6.350	104.768	-10.820	1.00	27.67	T6
	ATOM	5743	OD1	ASP	A	4	6.599	105.992	-10.846	1.00	30.20	T6
20	ATOM	5744	OD2	ASP	A	4	5.428	104.243	-11.489	1.00	29.12	T6
	ATOM	5745	C	ASP	A	4	9.223	103.669	-8.499	1.00	29.60	T6
	ATOM	5746	O	ASP	A	4	10.021	102.818	-8.884	1.00	34.27	T6
	ATOM	5747	N	CYS	A	5	8.948	103.880	-7.217	1.00	26.17	T6
	ATOM	5748	CA	CYS	A	5	9.550	103.092	-6.148	1.00	33.94	T6
25	ATOM	5749	CB	CYS	A	5	10.965	103.601	-5.807	1.00	34.14	T6
	ATOM	5750	SG	CYS	A	5	11.261	105.377	-6.009	1.00	28.49	T6
	ATOM	5751	C	CYS	A	5	8.666	103.135	-4.911	1.00	32.82	T6
	ATOM	5752	O	CYS	A	5	7.990	104.122	-4.658	1.00	31.80	T6
	ATOM	5753	N	LEU	A	6	8.647	102.042	-4.157	1.00	39.99	T6
30	ATOM	5754	CA	LEU	A	6	7.862	101.973	-2.930	1.00	33.84	T6
	ATOM	5755	CB	LEU	A	6	6.569	101.200	-3.162	1.00	29.40	T6
	ATOM	5756	CG	LEU	A	6	5.665	100.985	-1.949	1.00	23.57	T6
	ATOM	5757	CD1	LEU	A	6	4.233	100.877	-2.409	1.00	30.94	T6
	ATOM	5758	CD2	LEU	A	6	6.076	99.733	-1.207	1.00	32.12	T6
35	ATOM	5759	C	LEU	A	6	8.719	101.275	-1.894	1.00	32.24	T6
	ATOM	5760	O	LEU	A	6	9.339	100.263	-2.188	1.00	32.61	T6
	ATOM	5761	N	GLN	A	7	8.764	101.824	-0.686	1.00	28.14	T6
	ATOM	5762	CA	GLN	A	7	9.578	101.247	0.373	1.00	27.97	T6
	ATOM	5763	CB	GLN	A	7	10.787	102.137	0.638	1.00	34.28	T6
40	ATOM	5764	CG	GLN	A	7	11.841	101.507	1.516	1.00	29.43	T6
	ATOM	5765	CD	GLN	A	7	13.096	102.328	1.540	1.00	34.58	T6
	ATOM	5766	OE1	GLN	A	7	13.120	103.417	2.104	1.00	27.51	T6
	ATOM	5767	NE2	GLN	A	7	14.147	101.822	0.912	1.00	35.61	T6
	ATOM	5768	C	GLN	A	7	8.809	101.054	1.664	1.00	28.22	T6
45	ATOM	5769	O	GLN	A	7	8.011	101.904	2.060	1.00	26.49	T6
	ATOM	5770	N	LEU	A	8	9.065	99.932	2.323	1.00	32.35	T6
	ATOM	5771	CA	LEU	A	8	8.400	99.612	3.574	1.00	26.16	T6
	ATOM	5772	CB	LEU	A	8	7.675	98.270	3.449	1.00	29.17	T6
	ATOM	5773	CG	LEU	A	8	6.253	98.197	2.876	1.00	29.51	T6
50	ATOM	5774	CD1	LEU	A	8	5.877	99.478	2.185	1.00	29.61	T6
	ATOM	5775	CD2	LEU	A	8	6.167	97.021	1.933	1.00	24.52	T6
	ATOM	5776	C	LEU	A	8	9.380	99.566	4.740	1.00	36.10	T6
	ATOM	5777	O	LEU	A	8	10.564	99.313	4.562	1.00	30.42	T6
	ATOM	5778	N	ILE	A	9	8.864	99.810	5.937	1.00	34.53	T6
55	ATOM	5779	CA	ILE	A	9	9.649	99.815	7.163	1.00	36.78	T6
	ATOM	5780	CB	ILE	A	9	9.724	101.229	7.732	1.00	32.76	T6
	ATOM	5781	CG2	ILE	A	9	10.099	101.195	9.181	1.00	26.88	T6
	ATOM	5782	CG1	ILE	A	9	10.736	102.055	6.974	1.00	35.74	T6
	ATOM	5783	CD1	ILE	A	9	10.792	103.459	7.507	1.00	30.61	T6
60	ATOM	5784	C	ILE	A	9	8.962	98.941	8.208	1.00	32.82	T6
	ATOM	5785	O	ILE	A	9	7.741	98.904	8.277	1.00	32.53	T6
	ATOM	5786	N	ALA	A	10	9.736	98.253	9.036	1.00	28.41	T6
	ATOM	5787	CA	ALA	A	10	9.144	97.415	10.067	1.00	26.62	T6
	ATOM	5788	CB	ALA	A	10	10.216	96.678	10.828	1.00	29.09	T6
65	ATOM	5789	C	ALA	A	10	8.318	98.255	11.037	1.00	29.03	T6
	ATOM	5790	O	ALA	A	10	8.776	99.291	11.531	1.00	31.31	T6

	ATOM	5791	N	ASP	A	11	7.093	97.806	11.304	1.00	34.73	T6
	ATOM	5792	CA	ASP	A	11	6.210	98.499	12.230	1.00	24.60	T6
	ATOM	5793	CB	ASP	A	11	4.753	98.361	11.803	1.00	35.18	T6
	ATOM	5794	CG	ASP	A	11	3.805	98.961	12.817	1.00	27.93	T6
5	ATOM	5795	OD1	ASP	A	11	4.151	100.030	13.373	1.00	26.64	T6
	ATOM	5796	OD2	ASP	A	11	2.724	98.377	13.056	1.00	37.17	T6
	ATOM	5797	C	ASP	A	11	6.395	97.940	13.632	1.00	31.52	T6
	ATOM	5798	O	ASP	A	11	5.725	96.992	14.033	1.00	28.77	T6
	ATOM	5799	N	SER	A	12	7.319	98.559	14.362	1.00	32.20	T6
10	ATOM	5800	CA	SER	A	12	7.675	98.169	15.723	1.00	29.36	T6
	ATOM	5801	CB	SER	A	12	8.877	98.992	16.213	1.00	30.39	T6
	ATOM	5802	OG	SER	A	12	8.615	100.389	16.173	1.00	31.88	T6
	ATOM	5803	C	SER	A	12	6.540	98.324	16.713	1.00	31.92	T6
	ATOM	5804	O	SER	A	12	6.776	98.445	17.914	1.00	31.02	T6
15	ATOM	5805	N	GLU	A	13	5.306	98.326	16.228	1.00	35.05	T6
	ATOM	5806	CA	GLU	A	13	4.188	98.477	17.139	1.00	29.66	T6
	ATOM	5807	CB	GLU	A	13	3.730	99.922	17.157	1.00	33.13	T6
	ATOM	5808	CG	GLU	A	13	4.462	100.712	18.211	1.00	31.43	T6
	ATOM	5809	CD	GLU	A	13	3.930	102.123	18.316	1.00	31.06	T6
20	ATOM	5810	OE1	GLU	A	13	2.677	102.292	18.250	1.00	31.93	T6
	ATOM	5811	OE2	GLU	A	13	4.757	103.061	18.473	1.00	36.43	T6
	ATOM	5812	C	GLU	A	13	3.008	97.555	16.905	1.00	24.74	T6
	ATOM	5813	O	GLU	A	13	1.871	97.873	17.264	1.00	25.45	T6
	ATOM	5814	N	THR	A	14	3.284	96.414	16.290	1.00	31.90	T6
25	ATOM	5815	CA	THR	A	14	2.266	95.411	16.059	1.00	30.73	T6
	ATOM	5816	CB	THR	A	14	1.546	95.578	14.701	1.00	33.75	T6
	ATOM	5817	OG1	THR	A	14	2.505	95.550	13.646	1.00	32.99	T6
	ATOM	5818	CG2	THR	A	14	0.777	96.886	14.655	1.00	28.65	T6
	ATOM	5819	C	THR	A	14	3.013	94.095	16.091	1.00	32.63	T6
30	ATOM	5820	O	THR	A	14	4.199	94.031	15.769	1.00	28.71	T6
	ATOM	5821	N	PRO	A	15	2.330	93.026	16.503	1.00	36.36	T6
	ATOM	5822	CD	PRO	A	15	0.910	92.995	16.891	1.00	25.45	T6
	ATOM	5823	CA	PRO	A	15	2.928	91.692	16.591	1.00	25.43	T6
	ATOM	5824	CB	PRO	A	15	1.790	90.836	17.151	1.00	32.65	T6
35	ATOM	5825	CG	PRO	A	15	0.888	91.842	17.845	1.00	29.54	T6
	ATOM	5826	C	PRO	A	15	3.376	91.198	15.234	1.00	25.33	T6
	ATOM	5827	O	PRO	A	15	2.752	91.498	14.223	1.00	30.61	T6
	ATOM	5828	N	THR	A	16	4.457	90.439	15.212	1.00	24.17	T6
	ATOM	5829	CA	THR	A	16	4.943	89.900	13.952	1.00	27.65	T6
40	ATOM	5830	CB	THR	A	16	6.335	89.302	14.134	1.00	35.85	T6
	ATOM	5831	OG1	THR	A	16	6.251	88.109	14.927	1.00	37.63	T6
	ATOM	5832	CG2	THR	A	16	7.222	90.296	14.851	1.00	34.10	T6
	ATOM	5833	C	THR	A	16	3.982	88.804	13.501	1.00	32.73	T6
	ATOM	5834	O	THR	A	16	3.838	87.784	14.179	1.00	34.75	T6
45	ATOM	5835	N	ILE	A	17	3.330	89.009	12.362	1.00	30.16	T6
	ATOM	5836	CA	ILE	A	17	2.371	88.031	11.840	1.00	32.33	T6
	ATOM	5837	CB	ILE	A	17	1.875	88.453	10.459	1.00	33.67	T6
	ATOM	5838	CG2	ILE	A	17	0.898	87.435	9.931	1.00	29.51	T6
	ATOM	5839	CG1	ILE	A	17	1.213	89.826	10.552	1.00	27.87	T6
50	ATOM	5840	CD1	ILE	A	17	0.748	90.367	9.221	1.00	31.65	T6
	ATOM	5841	C	ILE	A	17	2.882	86.583	11.746	1.00	25.86	T6
	ATOM	5842	O	ILE	A	17	3.968	86.317	11.218	1.00	31.36	T6
	ATOM	5843	N	GLN	A	18	2.081	85.654	12.267	1.00	30.43	T6
	ATOM	5844	CA	GLN	A	18	2.414	84.233	12.242	1.00	31.16	T6
55	ATOM	5845	CB	GLN	A	18	2.368	83.644	13.643	1.00	37.60	T6
	ATOM	5846	CG	GLN	A	18	3.653	82.979	14.025	1.00	28.28	T6
	ATOM	5847	CD	GLN	A	18	4.722	83.993	14.310	1.00	35.84	T6
	ATOM	5848	OE1	GLN	A	18	4.723	84.627	15.368	1.00	25.77	T6
	ATOM	5849	NE2	GLN	A	18	5.636	84.178	13.357	1.00	24.73	T6
60	ATOM	5850	C	GLN	A	18	1.448	83.458	11.344	1.00	32.86	T6
	ATOM	5851	O	GLN	A	18	0.227	83.647	11.412	1.00	28.64	T6
	ATOM	5852	N	LYS	A	19	1.989	82.572	10.513	1.00	34.54	T6
	ATOM	5853	CA	LYS	A	19	1.148	81.807	9.599	1.00	30.11	T6
	ATOM	5854	CB	LYS	A	19	0.483	82.754	8.597	1.00	24.94	T6
65	ATOM	5855	CG	LYS	A	19	-0.209	82.043	7.441	1.00	19.80	T6
	ATOM	5856	CD	LYS	A	19	-1.032	83.032	6.614	1.00	33.17	T6

	ATOM	5857	CE	LYS	A	19	-1.730	82.336	5.450	1.00	39.36	T6
	ATOM	5858	NZ	LYS	A	19	-2.583	83.274	4.642	1.00	30.18	T6
	ATOM	5859	C	LYS	A	19	1.913	80.735	8.839	1.00	28.90	T6
	ATOM	5860	O	LYS	A	19	2.954	81.008	8.242	1.00	31.06	T6
5	ATOM	5861	N	GLY	A	20	1.372	79.521	8.845	1.00	29.72	T6
	ATOM	5862	CA	GLY	A	20	2.021	78.411	8.165	1.00	28.83	T6
	ATOM	5863	C	GLY	A	20	3.444	78.246	8.666	1.00	34.29	T6
	ATOM	5864	O	GLY	A	20	4.353	77.981	7.872	1.00	27.70	T6
	ATOM	5865	N	SER	A	21	3.634	78.404	9.979	1.00	26.99	T6
10	ATOM	5866	CA	SER	A	21	4.959	78.306	10.613	1.00	37.07	T6
	ATOM	5867	CB	SER	A	21	5.428	76.843	10.665	1.00	36.54	T6
	ATOM	5868	OG	SER	A	21	5.571	76.293	9.368	1.00	29.00	T6
	ATOM	5869	C	SER	A	21	6.021	79.190	9.921	1.00	29.35	T6
	ATOM	5870	O	SER	A	21	7.184	78.802	9.757	1.00	35.93	T6
15	ATOM	5871	N	TYR	A	22	5.584	80.383	9.527	1.00	35.00	T6
	ATOM	5872	CA	TYR	A	22	6.418	81.380	8.874	1.00	29.65	T6
	ATOM	5873	CB	TYR	A	22	5.968	81.583	7.432	1.00	33.78	T6
	ATOM	5874	CG	TYR	A	22	6.738	80.783	6.423	1.00	29.35	T6
	ATOM	5875	CD1	TYR	A	22	7.374	79.597	6.782	1.00	22.46	T6
20	ATOM	5876	CE1	TYR	A	22	8.066	78.834	5.839	1.00	28.92	T6
	ATOM	5877	CD2	TYR	A	22	6.809	81.193	5.093	1.00	26.90	T6
	ATOM	5878	CE2	TYR	A	22	7.496	80.437	4.138	1.00	32.78	T6
	ATOM	5879	CZ	TYR	A	22	8.121	79.258	4.523	1.00	30.31	T6
	ATOM	5880	OH	TYR	A	22	8.796	78.497	3.594	1.00	25.02	T6
25	ATOM	5881	C	TYR	A	22	6.197	82.673	9.638	1.00	32.38	T6
	ATOM	5882	O	TYR	A	22	5.098	82.912	10.152	1.00	31.53	T6
	ATOM	5883	N	THR	A	23	7.231	83.503	9.728	1.00	29.83	T6
	ATOM	5884	CA	THR	A	23	7.081	84.775	10.421	1.00	30.64	T6
	ATOM	5885	CB	THR	A	23	8.219	85.032	11.424	1.00	33.52	T6
30	ATOM	5886	OG1	THR	A	23	8.541	83.825	12.121	1.00	26.48	T6
	ATOM	5887	CG2	THR	A	23	7.777	86.058	12.438	1.00	38.89	T6
	ATOM	5888	C	THR	A	23	7.078	85.886	9.378	1.00	30.50	T6
	ATOM	5889	O	THR	A	23	7.932	85.926	8.489	1.00	33.17	T6
	ATOM	5890	N	PHE	A	24	6.095	86.772	9.476	1.00	30.85	T6
35	ATOM	5891	CA	PHE	A	24	5.990	87.885	8.546	1.00	27.46	T6
	ATOM	5892	CB	PHE	A	24	4.669	87.814	7.789	1.00	31.32	T6
	ATOM	5893	CG	PHE	A	24	4.560	86.630	6.881	1.00	24.94	T6
	ATOM	5894	CD1	PHE	A	24	4.214	85.383	7.378	1.00	36.74	T6
	ATOM	5895	CD2	PHE	A	24	4.827	86.756	5.521	1.00	28.85	T6
40	ATOM	5896	CE1	PHE	A	24	4.134	84.275	6.531	1.00	30.39	T6
	ATOM	5897	CE2	PHE	A	24	4.751	85.658	4.669	1.00	29.62	T6
	ATOM	5898	CZ	PHE	A	24	4.404	84.415	5.178	1.00	31.64	T6
	ATOM	5899	C	PHE	A	24	6.108	89.235	9.256	1.00	31.74	T6
	ATOM	5900	O	PHE	A	24	5.362	89.538	10.190	1.00	31.62	T6
45	ATOM	5901	N	VAL	A	25	7.060	90.041	8.802	1.00	30.77	T6
	ATOM	5902	CA	VAL	A	25	7.289	91.357	9.375	1.00	28.29	T6
	ATOM	5903	CB	VAL	A	25	8.527	92.020	8.756	1.00	31.62	T6
	ATOM	5904	CG1	VAL	A	25	8.701	93.417	9.303	1.00	28.92	T6
	ATOM	5905	CG2	VAL	A	25	9.751	91.185	9.042	1.00	31.05	T6
50	ATOM	5906	C	VAL	A	25	6.102	92.271	9.124	1.00	29.12	T6
	ATOM	5907	O	VAL	A	25	5.585	92.336	8.012	1.00	30.56	T6
	ATOM	5908	N	PRO	A	26	5.644	92.982	10.166	1.00	31.19	T6
	ATOM	5909	CD	PRO	A	26	6.054	92.823	11.567	1.00	32.73	T6
	ATOM	5910	CA	PRO	A	26	4.511	93.910	10.062	1.00	30.19	T6
55	ATOM	5911	CB	PRO	A	26	4.158	94.197	11.516	1.00	32.57	T6
	ATOM	5912	CG	PRO	A	26	4.770	93.057	12.276	1.00	31.54	T6
	ATOM	5913	C	PRO	A	26	5.035	95.165	9.372	1.00	27.56	T6
	ATOM	5914	O	PRO	A	26	5.836	95.891	9.954	1.00	22.67	T6
	ATOM	5915	N	TRP	A	27	4.601	95.431	8.146	1.00	31.50	T6
60	ATOM	5916	CA	TRP	A	27	5.100	96.605	7.440	1.00	28.90	T6
	ATOM	5917	CB	TRP	A	27	5.131	96.349	5.929	1.00	31.55	T6
	ATOM	5918	CG	TRP	A	27	6.057	95.264	5.543	1.00	25.84	T6
	ATOM	5919	CD2	TRP	A	27	7.447	95.162	5.867	1.00	28.59	T6
	ATOM	5920	CE2	TRP	A	27	7.908	93.934	5.348	1.00	26.93	T6
65	ATOM	5921	CE3	TRP	A	27	8.345	95.983	6.547	1.00	34.99	T6
	ATOM	5922	CD1	TRP	A	27	5.741	94.135	4.856	1.00	32.48	T6

	ATOM	5923	NE1	TRP	A	27	6.846	93.324	4.735	1.00	26.77	T6
	ATOM	5924	CZ2	TRP	A	27	9.234	93.509	5.488	1.00	29.65	T6
	ATOM	5925	CZ3	TRP	A	27	9.667	95.556	6.684	1.00	28.47	T6
	ATOM	5926	CH2	TRP	A	27	10.094	94.331	6.158	1.00	41.29	T6
5	ATOM	5927	C	TRP	A	27	4.371	97.918	7.718	1.00	28.09	T6
	ATOM	5928	O	TRP	A	27	3.268	97.953	8.260	1.00	40.32	T6
	ATOM	5929	N	LEU	A	28	5.036	99.000	7.334	1.00	31.39	T6
	ATOM	5930	CA	LEU	A	28	4.530	100.347	7.489	1.00	34.93	T6
	ATOM	5931	CB	LEU	A	28	4.971	100.908	8.829	1.00	30.47	T6
10	ATOM	5932	CG	LEU	A	28	4.154	102.093	9.324	1.00	34.73	T6
	ATOM	5933	CD1	LEU	A	28	2.715	101.637	9.560	1.00	34.96	T6
	ATOM	5934	CD2	LEU	A	28	4.775	102.636	10.601	1.00	33.59	T6
	ATOM	5935	C	LEU	A	28	5.145	101.153	6.351	1.00	32.22	T6
	ATOM	5936	O	LEU	A	28	6.358	101.129	6.146	1.00	33.85	T6
15	ATOM	5937	N	LEU	A	29	4.317	101.864	5.603	1.00	30.67	T6
	ATOM	5938	CA	LEU	A	29	4.826	102.627	4.480	1.00	28.44	T6
	ATOM	5939	CB	LEU	A	29	3.711	103.423	3.820	1.00	37.43	T6
	ATOM	5940	CG	LEU	A	29	4.206	104.265	2.648	1.00	28.27	T6
	ATOM	5941	CD1	LEU	A	29	4.452	103.365	1.457	1.00	38.03	T6
20	ATOM	5942	CD2	LEU	A	29	3.196	105.324	2.303	1.00	27.49	T6
	ATOM	5943	C	LEU	A	29	5.931	103.585	4.857	1.00	28.38	T6
	ATOM	5944	O	LEU	A	29	5.764	104.405	5.757	1.00	26.72	T6
	ATOM	5945	N	SER	A	30	7.064	103.471	4.168	1.00	31.21	T6
	ATOM	5946	CA	SER	A	30	8.180	104.379	4.393	1.00	29.16	T6
25	ATOM	5947	CB	SER	A	30	9.500	103.729	4.017	1.00	31.88	T6
	ATOM	5948	OG	SER	A	30	10.550	104.668	4.138	1.00	37.16	T6
	ATOM	5949	C	SER	A	30	7.920	105.548	3.454	1.00	26.62	T6
	ATOM	5950	O	SER	A	30	7.890	106.703	3.858	1.00	35.86	T6
	ATOM	5951	N	PHE	A	31	7.723	105.224	2.186	1.00	25.03	T6
30	ATOM	5952	CA	PHE	A	31	7.432	106.232	1.176	1.00	23.32	T6
	ATOM	5953	CB	PHE	A	31	8.662	107.107	0.920	1.00	24.44	T6
	ATOM	5954	CG	PHE	A	31	9.593	106.559	-0.118	1.00	33.57	T6
	ATOM	5955	CD1	PHE	A	31	9.355	106.780	-1.469	1.00	28.19	T6
	ATOM	5956	CD2	PHE	A	31	10.706	105.819	0.255	1.00	34.21	T6
35	ATOM	5957	CE1	PHE	A	31	10.214	106.272	-2.434	1.00	31.94	T6
	ATOM	5958	CE2	PHE	A	31	11.568	105.307	-0.697	1.00	29.18	T6
	ATOM	5959	CZ	PHE	A	31	11.324	105.533	-2.045	1.00	25.06	T6
	ATOM	5960	C	PHE	A	31	6.992	105.534	-0.115	1.00	25.83	T6
	ATOM	5961	O	PHE	A	31	7.367	104.394	-0.380	1.00	32.83	T6
40	ATOM	5962	N	LYS	A	32	6.180	106.221	-0.902	1.00	26.40	T6
	ATOM	5963	CA	LYS	A	32	5.694	105.672	-2.153	1.00	33.58	T6
	ATOM	5964	CB	LYS	A	32	4.235	105.251	-2.014	1.00	28.79	T6
	ATOM	5965	CG	LYS	A	32	3.592	104.952	-3.336	1.00	33.62	T6
	ATOM	5966	CD	LYS	A	32	2.111	104.750	-3.212	1.00	36.78	T6
45	ATOM	5967	CE	LYS	A	32	1.505	104.722	-4.596	1.00	29.00	T6
	ATOM	5968	NZ	LYS	A	32	0.069	104.351	-4.549	1.00	32.86	T6
	ATOM	5969	C	LYS	A	32	5.826	106.737	-3.230	1.00	28.08	T6
	ATOM	5970	O	LYS	A	32	5.344	107.856	-3.075	1.00	28.42	T6
	ATOM	5971	N	ARG	A	33	6.480	106.389	-4.325	1.00	31.03	T6
50	ATOM	5972	CA	ARG	A	33	6.697	107.336	-5.408	1.00	30.51	T6
	ATOM	5973	CB	ARG	A	33	8.186	107.694	-5.461	1.00	31.76	T6
	ATOM	5974	CG	ARG	A	33	8.638	108.543	-6.622	1.00	31.05	T6
	ATOM	5975	CD	ARG	A	33	9.949	109.217	-6.258	1.00	44.32	T6
	ATOM	5976	NE	ARG	A	33	10.531	109.989	-7.349	1.00	32.73	T6
55	ATOM	5977	CZ	ARG	A	33	11.245	109.455	-8.335	1.00	29.96	T6
	ATOM	5978	NH1	ARG	A	33	11.467	108.145	-8.356	1.00	35.60	T6
	ATOM	5979	NH2	ARG	A	33	11.717	110.226	-9.308	1.00	34.01	T6
	ATOM	5980	C	ARG	A	33	6.240	106.736	-6.726	1.00	26.24	T6
	ATOM	5981	O	ARG	A	33	6.695	105.661	-7.119	1.00	33.74	T6
60	ATOM	5982	N	GLY	A	34	5.330	107.424	-7.405	1.00	30.29	T6
	ATOM	5983	CA	GLY	A	34	4.849	106.917	-8.673	1.00	31.40	T6
	ATOM	5984	C	GLY	A	34	3.593	106.078	-8.555	1.00	32.25	T6
	ATOM	5985	O	GLY	A	34	2.919	106.074	-7.516	1.00	31.25	T6
	ATOM	5986	N	SER	A	35	3.289	105.344	-9.618	1.00	23.45	T6
65	ATOM	5987	CA	SER	A	35	2.085	104.529	-9.647	1.00	27.27	T6
	ATOM	5988	CB	SER	A	35	1.185	105.027	-10.771	1.00	33.95	T6

	ATOM	5989	OG	SER	A	35	1.901	105.014	-11.995	1.00	33.31	T6
	ATOM	5990	C	SER	A	35	2.290	103.028	-9.825	1.00	35.58	T6
	ATOM	5991	O	SER	A	35	1.404	102.243	-9.493	1.00	28.04	T6
	ATOM	5992	N	ALA	A	36	3.442	102.624	-10.348	1.00	35.08	T6
5	ATOM	5993	CA	ALA	A	36	3.707	101.210	-10.596	1.00	34.04	T6
	ATOM	5994	CB	ALA	A	36	5.029	101.063	-11.332	1.00	25.21	T6
	ATOM	5995	C	ALA	A	36	3.683	100.267	-9.393	1.00	29.16	T6
	ATOM	5996	O	ALA	A	36	3.586	99.056	-9.571	1.00	31.71	T6
	ATOM	5997	N	LEU	A	37	3.766	100.806	-8.179	1.00	29.34	T6
10	ATOM	5998	CA	LEU	A	37	3.768	99.967	-6.978	1.00	24.72	T6
	ATOM	5999	CB	LEU	A	37	5.198	99.832	-6.454	1.00	33.68	T6
	ATOM	6000	CG	LEU	A	37	6.174	99.190	-7.441	1.00	28.94	T6
	ATOM	6001	CD1	LEU	A	37	7.600	99.494	-7.045	1.00	29.68	T6
	ATOM	6002	CD2	LEU	A	37	5.933	97.706	-7.494	1.00	29.50	T6
15	ATOM	6003	C	LEU	A	37	2.854	100.489	-5.862	1.00	24.02	T6
	ATOM	6004	O	LEU	A	37	2.693	101.696	-5.690	1.00	37.30	T6
	ATOM	6005	N	GLU	A	38	2.267	99.567	-5.103	1.00	32.19	T6
	ATOM	6006	CA	GLU	A	38	1.351	99.915	-4.016	1.00	33.60	T6
	ATOM	6007	CB	GLU	A	38	-0.091	99.871	-4.515	1.00	27.16	T6
20	ATOM	6008	CG	GLU	A	38	-0.580	101.105	-5.246	1.00	34.43	T6
	ATOM	6009	CD	GLU	A	38	-1.919	100.860	-5.946	1.00	28.89	T6
	ATOM	6010	OE1	GLU	A	38	-2.764	100.115	-5.377	1.00	29.78	T6
	ATOM	6011	OE2	GLU	A	38	-2.123	101.418	-7.059	1.00	30.19	T6
	ATOM	6012	C	GLU	A	38	1.453	98.951	-2.842	1.00	25.83	T6
25	ATOM	6013	O	GLU	A	38	2.001	97.857	-2.976	1.00	28.89	T6
	ATOM	6014	N	GLU	A	39	0.920	99.357	-1.689	1.00	32.89	T6
	ATOM	6015	CA	GLU	A	39	0.904	98.482	-0.519	1.00	29.88	T6
	ATOM	6016	CB	GLU	A	39	0.871	99.247	0.787	1.00	36.44	T6
	ATOM	6017	CG	GLU	A	39	1.758	100.418	0.873	1.00	31.24	T6
30	ATOM	6018	CD	GLU	A	39	1.172	101.454	1.811	1.00	34.86	T6
	ATOM	6019	OE1	GLU	A	39	0.512	102.403	1.311	1.00	27.40	T6
	ATOM	6020	OE2	GLU	A	39	1.346	101.303	3.048	1.00	28.17	T6
	ATOM	6021	C	GLU	A	39	-0.423	97.753	-0.586	1.00	26.97	T6
	ATOM	6022	O	GLU	A	39	-1.430	98.318	-1.019	1.00	31.22	T6
35	ATOM	6023	N	LYS	A	40	-0.440	96.510	-0.142	1.00	30.65	T6
	ATOM	6024	CA	LYS	A	40	-1.678	95.765	-0.145	1.00	28.24	T6
	ATOM	6025	CB	LYS	A	40	-2.012	95.252	-1.543	1.00	28.86	T6
	ATOM	6026	CG	LYS	A	40	-3.344	94.503	-1.590	1.00	31.27	T6
	ATOM	6027	CD	LYS	A	40	-3.484	93.694	-2.864	1.00	32.57	T6
40	ATOM	6028	CE	LYS	A	40	-4.702	92.782	-2.804	1.00	30.86	T6
	ATOM	6029	NZ	LYS	A	40	-4.745	91.882	-3.999	1.00	30.46	T6
	ATOM	6030	C	LYS	A	40	-1.583	94.601	0.810	1.00	26.29	T6
	ATOM	6031	O	LYS	A	40	-0.946	93.589	0.519	1.00	27.96	T6
	ATOM	6032	N	GLU	A	41	-2.208	94.761	1.967	1.00	27.85	T6
45	ATOM	6033	CA	GLU	A	41	-2.215	93.718	2.970	1.00	30.65	T6
	ATOM	6034	CB	GLU	A	41	-3.133	92.598	2.501	1.00	31.31	T6
	ATOM	6035	CG	GLU	A	41	-4.491	93.139	2.073	1.00	31.69	T6
	ATOM	6036	CD	GLU	A	41	-5.313	92.141	1.257	1.00	30.49	T6
	ATOM	6037	OE1	GLU	A	41	-4.790	91.610	0.234	1.00	30.53	T6
50	ATOM	6038	OE2	GLU	A	41	-6.494	91.899	1.631	1.00	27.73	T6
	ATOM	6039	C	GLU	A	41	-0.810	93.195	3.250	1.00	34.93	T6
	ATOM	6040	O	GLU	A	41	-0.516	92.010	3.072	1.00	26.65	T6
	ATOM	6041	N	ASN	A	42	0.058	94.104	3.671	1.00	31.06	T6
	ATOM	6042	CA	ASN	A	42	1.425	93.765	4.024	1.00	27.05	T6
55	ATOM	6043	CB	ASN	A	42	1.425	92.781	5.179	1.00	35.52	T6
	ATOM	6044	CG	ASN	A	42	2.454	93.122	6.204	1.00	29.68	T6
	ATOM	6045	OD1	ASN	A	42	3.279	92.292	6.567	1.00	31.70	T6
	ATOM	6046	ND2	ASN	A	42	2.421	94.358	6.684	1.00	25.07	T6
	ATOM	6047	C	ASN	A	42	2.277	93.212	2.899	1.00	34.53	T6
60	ATOM	6048	O	ASN	A	42	3.289	92.565	3.149	1.00	28.81	T6
	ATOM	6049	N	LYS	A	43	1.870	93.467	1.662	1.00	32.81	T6
	ATOM	6050	CA	LYS	A	43	2.618	93.004	0.504	1.00	39.82	T6
	ATOM	6051	CB	LYS	A	43	1.907	91.830	-0.157	1.00	31.56	T6
	ATOM	6052	CG	LYS	A	43	1.922	90.560	0.664	1.00	35.73	T6
65	ATOM	6053	CD	LYS	A	43	1.133	89.449	-0.009	1.00	35.13	T6
	ATOM	6054	CE	LYS	A	43	-0.360	89.731	0.043	1.00	25.26	T6

	ATOM	6055	NZ	LYS	A	43	-1.163	88.636	-0.581	1.00	25.44	T6
	ATOM	6056	C	LYS	A	43	2.731	94.143	-0.486	1.00	36.22	T6
	ATOM	6057	O	LYS	A	43	1.980	95.107	-0.410	1.00	23.99	T6
	ATOM	6058	N	ILE	A	44	3.677	94.046	-1.408	1.00	35.30	T6
5	ATOM	6059	CA	ILE	A	44	3.835	95.084	-2.408	1.00	32.65	T6
	ATOM	6060	CB	ILE	A	44	5.304	95.330	-2.731	1.00	21.78	T6
	ATOM	6061	CG2	ILE	A	44	5.422	96.403	-3.790	1.00	30.35	T6
	ATOM	6062	CG1	ILE	A	44	6.043	95.753	-1.466	1.00	28.84	T6
	ATOM	6063	CD1	ILE	A	44	7.504	96.003	-1.665	1.00	27.10	T6
10	ATOM	6064	C	ILE	A	44	3.125	94.618	-3.661	1.00	35.35	T6
	ATOM	6065	O	ILE	A	44	3.401	93.536	-4.175	1.00	34.17	T6
	ATOM	6066	N	LEU	A	45	2.197	95.433	-4.143	1.00	29.15	T6
	ATOM	6067	CA	LEU	A	45	1.438	95.088	-5.332	1.00	25.96	T6
	ATOM	6068	CB	LEU	A	45	-0.027	95.473	-5.131	1.00	24.19	T6
15	ATOM	6069	CG	LEU	A	45	-0.915	95.247	-6.363	1.00	27.00	T6
	ATOM	6070	CD1	LEU	A	45	-1.120	93.751	-6.575	1.00	30.10	T6
	ATOM	6071	CD2	LEU	A	45	-2.234	95.954	-6.171	1.00	25.80	T6
	ATOM	6072	C	LEU	A	45	1.962	95.747	-6.611	1.00	24.61	T6
	ATOM	6073	O	LEU	A	45	2.143	96.967	-6.670	1.00	25.73	T6
20	ATOM	6074	N	VAL	A	46	2.183	94.941	-7.642	1.00	28.45	T6
	ATOM	6075	CA	VAL	A	46	2.677	95.460	-8.909	1.00	30.26	T6
	ATOM	6076	CB	VAL	A	46	3.428	94.370	-9.680	1.00	33.66	T6
	ATOM	6077	CG1	VAL	A	46	3.935	94.924	-10.992	1.00	25.07	T6
	ATOM	6078	CG2	VAL	A	46	4.574	93.861	-8.853	1.00	36.90	T6
25	ATOM	6079	C	VAL	A	46	1.532	95.983	-9.780	1.00	25.05	T6
	ATOM	6080	O	VAL	A	46	0.631	95.228	-10.147	1.00	28.49	T6
	ATOM	6081	N	LYS	A	47	1.571	97.269	-10.122	1.00	33.33	T6
	ATOM	6082	CA	LYS	A	47	0.517	97.860	-10.941	1.00	36.52	T6
	ATOM	6083	CB	LYS	A	47	0.091	99.208	-10.362	1.00	29.80	T6
30	ATOM	6084	CG	LYS	A	47	-0.777	99.107	-9.122	1.00	32.43	T6
	ATOM	6085	CD	LYS	A	47	-1.977	98.229	-9.406	1.00	26.80	T6
	ATOM	6086	CE	LYS	A	47	-3.028	98.334	-8.313	1.00	37.19	T6
	ATOM	6087	NZ	LYS	A	47	-3.772	99.630	-8.370	1.00	32.14	T6
	ATOM	6088	C	LYS	A	47	0.868	98.040	-12.410	1.00	24.90	T6
35	ATOM	6089	O	LYS	A	47	0.005	98.375	-13.212	1.00	31.08	T6
	ATOM	6090	N	GLU	A	48	2.129	97.826	-12.762	1.00	37.36	T6
	ATOM	6091	CA	GLU	A	48	2.584	97.961	-14.141	1.00	32.36	T6
	ATOM	6092	CB	GLU	A	48	3.231	99.313	-14.370	1.00	24.60	T6
	ATOM	6093	CG	GLU	A	48	2.298	100.481	-14.356	1.00	34.25	T6
40	ATOM	6094	CD	GLU	A	48	3.048	101.794	-14.446	1.00	28.00	T6
	ATOM	6095	OE1	GLU	A	48	4.012	101.869	-15.248	1.00	31.33	T6
	ATOM	6096	OE2	GLU	A	48	2.669	102.746	-13.715	1.00	32.63	T6
	ATOM	6097	C	GLU	A	48	3.639	96.918	-14.360	1.00	33.94	T6
	ATOM	6098	O	GLU	A	48	4.611	96.871	-13.611	1.00	31.58	T6
45	ATOM	6099	N	THR	A	49	3.482	96.088	-15.382	1.00	24.51	T6
	ATOM	6100	CA	THR	A	49	4.488	95.063	-15.614	1.00	24.51	T6
	ATOM	6101	CB	THR	A	49	4.019	94.038	-16.643	1.00	31.41	T6
	ATOM	6102	OG1	THR	A	49	4.423	94.462	-17.940	1.00	27.65	T6
	ATOM	6103	CG2	THR	A	49	2.508	93.911	-16.613	1.00	31.06	T6
50	ATOM	6104	C	THR	A	49	5.782	95.722	-16.093	1.00	31.16	T6
	ATOM	6105	O	THR	A	49	5.758	96.775	-16.726	1.00	27.30	T6
	ATOM	6106	N	GLY	A	50	6.910	95.105	-15.761	1.00	30.92	T6
	ATOM	6107	CA	GLY	A	50	8.197	95.641	-16.160	1.00	36.81	T6
	ATOM	6108	C	GLY	A	50	9.307	94.994	-15.362	1.00	26.83	T6
55	ATOM	6109	O	GLY	A	50	9.092	93.958	-14.732	1.00	32.34	T6
	ATOM	6110	N	TYR	A	51	10.493	95.597	-15.391	1.00	26.17	T6
	ATOM	6111	CA	TYR	A	51	11.643	95.085	-14.647	1.00	30.59	T6
	ATOM	6112	CB	TYR	A	51	12.902	95.143	-15.511	1.00	30.53	T6
	ATOM	6113	CG	TYR	A	51	12.914	94.079	-16.570	1.00	35.13	T6
60	ATOM	6114	CD1	TYR	A	51	12.186	94.228	-17.744	1.00	25.41	T6
	ATOM	6115	CE1	TYR	A	51	12.099	93.187	-18.671	1.00	28.24	T6
	ATOM	6116	CD2	TYR	A	51	13.569	92.868	-16.349	1.00	35.87	T6
	ATOM	6117	CE2	TYR	A	51	13.490	91.822	-17.262	1.00	33.86	T6
	ATOM	6118	CZ	TYR	A	51	12.750	91.980	-18.418	1.00	29.46	T6
65	ATOM	6119	OH	TYR	A	51	12.620	90.908	-19.281	1.00	38.63	T6
	ATOM	6120	C	TYR	A	51	11.846	95.875	-13.357	1.00	29.08	T6

	ATOM	6121	O	TYR	A	51	11.908	97.105	-13.368	1.00	27.63	T6
	ATOM	6122	N	PHE	A	52	11.944	95.166	-12.241	1.00	31.10	T6
	ATOM	6123	CA	PHE	A	52	12.120	95.827	-10.961	1.00	29.55	T6
	ATOM	6124	CB	PHE	A	52	10.886	95.618	-10.081	1.00	25.19	T6
5	ATOM	6125	CG	PHE	A	52	9.605	96.111	-10.687	1.00	30.53	T6
	ATOM	6126	CD1	PHE	A	52	8.984	95.403	-11.707	1.00	20.87	T6
	ATOM	6127	CD2	PHE	A	52	9.009	97.280	-10.219	1.00	31.65	T6
	ATOM	6128	CE1	PHE	A	52	7.787	95.849	-12.251	1.00	28.78	T6
	ATOM	6129	CE2	PHE	A	52	7.817	97.735	-10.753	1.00	29.74	T6
10	ATOM	6130	CZ	PHE	A	52	7.201	97.020	-11.772	1.00	28.00	T6
	ATOM	6131	C	PHE	A	52	13.334	95.359	-10.179	1.00	28.20	T6
	ATOM	6132	O	PHE	A	52	13.745	94.209	-10.273	1.00	30.05	T6
	ATOM	6133	N	PHE	A	53	13.901	96.280	-9.409	1.00	38.46	T6
	ATOM	6134	CA	PHE	A	53	15.032	95.991	-8.548	1.00	30.31	T6
15	ATOM	6135	CB	PHE	A	53	15.979	97.180	-8.479	1.00	31.86	T6
	ATOM	6136	CG	PHE	A	53	17.052	97.037	-7.444	1.00	34.13	T6
	ATOM	6137	CD1	PHE	A	53	18.053	96.092	-7.584	1.00	33.67	T6
	ATOM	6138	CD2	PHE	A	53	17.051	97.842	-6.314	1.00	25.29	T6
	ATOM	6139	CE1	PHE	A	53	19.032	95.950	-6.616	1.00	35.39	T6
20	ATOM	6140	CE2	PHE	A	53	18.029	97.707	-5.337	1.00	40.12	T6
	ATOM	6141	CZ	PHE	A	53	19.017	96.762	-5.490	1.00	31.80	T6
	ATOM	6142	C	PHE	A	53	14.355	95.810	-7.209	1.00	31.32	T6
	ATOM	6143	O	PHE	A	53	13.645	96.703	-6.760	1.00	26.78	T6
	ATOM	6144	N	ILE	A	54	14.561	94.665	-6.571	1.00	28.52	T6
25	ATOM	6145	CA	ILE	A	54	13.914	94.394	-5.295	1.00	29.10	T6
	ATOM	6146	CB	ILE	A	54	12.982	93.190	-5.429	1.00	29.98	T6
	ATOM	6147	CG2	ILE	A	54	12.153	93.035	-4.179	1.00	27.16	T6
	ATOM	6148	CG1	ILE	A	54	12.086	93.386	-6.647	1.00	29.52	T6
	ATOM	6149	CD1	ILE	A	54	11.477	92.132	-7.157	1.00	29.78	T6
30	ATOM	6150	C	ILE	A	54	14.937	94.122	-4.210	1.00	27.60	T6
	ATOM	6151	O	ILE	A	54	15.849	93.319	-4.399	1.00	33.41	T6
	ATOM	6152	N	TYR	A	55	14.778	94.786	-3.071	1.00	28.61	T6
	ATOM	6153	CA	TYR	A	55	15.710	94.623	-1.967	1.00	29.34	T6
	ATOM	6154	CB	TYR	A	55	16.619	95.851	-1.878	1.00	39.51	T6
35	ATOM	6155	CG	TYR	A	55	15.891	97.160	-1.712	1.00	30.69	T6
	ATOM	6156	CD1	TYR	A	55	15.689	97.713	-0.456	1.00	28.08	T6
	ATOM	6157	CE1	TYR	A	55	14.985	98.892	-0.301	1.00	31.69	T6
	ATOM	6158	CD2	TYR	A	55	15.374	97.826	-2.815	1.00	29.32	T6
	ATOM	6159	CE2	TYR	A	55	14.666	99.007	-2.673	1.00	29.49	T6
40	ATOM	6160	CZ	TYR	A	55	14.471	99.537	-1.412	1.00	30.12	T6
	ATOM	6161	OH	TYR	A	55	13.748	100.698	-1.259	1.00	28.19	T6
	ATOM	6162	C	TYR	A	55	15.011	94.394	-0.644	1.00	28.56	T6
	ATOM	6163	O	TYR	A	55	13.800	94.499	-0.553	1.00	37.67	T6
	ATOM	6164	N	GLY	A	56	15.781	94.067	0.383	1.00	33.06	T6
45	ATOM	6165	CA	GLY	A	56	15.193	93.835	1.682	1.00	34.61	T6
	ATOM	6166	C	GLY	A	56	16.228	93.413	2.693	1.00	28.86	T6
	ATOM	6167	O	GLY	A	56	17.118	92.636	2.371	1.00	26.29	T6
	ATOM	6168	N	GLN	A	57	16.111	93.940	3.911	1.00	28.18	T6
	ATOM	6169	CA	GLN	A	57	17.021	93.620	5.007	1.00	29.05	T6
50	ATOM	6170	CB	GLN	A	57	17.993	94.780	5.267	1.00	30.47	T6
	ATOM	6171	CG	GLN	A	57	18.869	94.576	6.495	1.00	35.60	T6
	ATOM	6172	CD	GLN	A	57	19.918	95.654	6.683	1.00	34.53	T6
	ATOM	6173	OE1	GLN	A	57	20.869	95.751	5.914	1.00	34.52	T6
	ATOM	6174	NE2	GLN	A	57	19.752	96.467	7.719	1.00	26.12	T6
55	ATOM	6175	C	GLN	A	57	16.234	93.326	6.277	1.00	30.86	T6
	ATOM	6176	O	GLN	A	57	15.168	93.887	6.499	1.00	29.09	T6
	ATOM	6177	N	VAL	A	58	16.772	92.440	7.104	1.00	37.79	T6
	ATOM	6178	CA	VAL	A	58	16.152	92.060	8.369	1.00	27.95	T6
	ATOM	6179	CB	VAL	A	58	15.408	90.710	8.247	1.00	32.47	T6
60	ATOM	6180	CG1	VAL	A	58	15.011	90.198	9.619	1.00	34.50	T6
	ATOM	6181	CG2	VAL	A	58	14.189	90.874	7.381	1.00	29.70	T6
	ATOM	6182	C	VAL	A	58	17.249	91.905	9.416	1.00	34.21	T6
	ATOM	6183	O	VAL	A	58	18.318	91.401	9.112	1.00	28.26	T6
	ATOM	6184	N	LEU	A	59	16.992	92.346	10.640	1.00	30.12	T6
65	ATOM	6185	CA	LEU	A	59	17.974	92.210	11.710	1.00	34.19	T6
	ATOM	6186	CB	LEU	A	59	18.012	93.470	12.581	1.00	32.07	T6

	ATOM	6187	CG	LEU	A	59	19.305	93.741	13.363	1.00	33.38	T6
	ATOM	6188	CD1	LEU	A	59	19.027	94.734	14.456	1.00	26.27	T6
	ATOM	6189	CD2	LEU	A	59	19.855	92.472	13.970	1.00	28.05	T6
	ATOM	6190	C	LEU	A	59	17.568	91.017	12.577	1.00	33.35	T6
5	ATOM	6191	O	LEU	A	59	16.530	91.056	13.247	1.00	31.38	T6
	ATOM	6192	N	TYR	A	60	18.378	89.961	12.570	1.00	29.96	T6
	ATOM	6193	CA	TYR	A	60	18.068	88.778	13.368	1.00	28.88	T6
	ATOM	6194	CB	TYR	A	60	18.536	87.528	12.643	1.00	35.18	T6
	ATOM	6195	CG	TYR	A	60	17.880	87.389	11.318	1.00	35.29	T6
10	ATOM	6196	CD1	TYR	A	60	18.574	87.647	10.149	1.00	28.29	T6
	ATOM	6197	CE1	TYR	A	60	17.943	87.583	8.911	1.00	33.40	T6
	ATOM	6198	CD2	TYR	A	60	16.539	87.057	11.231	1.00	29.73	T6
	ATOM	6199	CE2	TYR	A	60	15.895	86.990	10.013	1.00	34.33	T6
	ATOM	6200	CZ	TYR	A	60	16.602	87.257	8.854	1.00	28.16	T6
15	ATOM	6201	OH	TYR	A	60	15.958	87.222	7.641	1.00	35.66	T6
	ATOM	6202	C	TYR	A	60	18.665	88.794	14.770	1.00	27.49	T6
	ATOM	6203	O	TYR	A	60	19.872	88.971	14.947	1.00	31.37	T6
	ATOM	6204	N	THR	A	61	17.814	88.601	15.765	1.00	30.01	T6
	ATOM	6205	CA	THR	A	61	18.262	88.582	17.142	1.00	32.89	T6
20	ATOM	6206	CB	THR	A	61	17.592	89.696	17.942	1.00	31.35	T6
	ATOM	6207	OG1	THR	A	61	16.171	89.604	17.788	1.00	30.49	T6
	ATOM	6208	CG2	THR	A	61	18.067	91.051	17.450	1.00	31.66	T6
	ATOM	6209	C	THR	A	61	17.908	87.230	17.733	1.00	28.20	T6
	ATOM	6210	O	THR	A	61	17.905	87.040	18.946	1.00	29.96	T6
25	ATOM	6211	N	ASP	A	62	17.613	86.290	16.846	1.00	29.48	T6
	ATOM	6212	CA	ASP	A	62	17.260	84.929	17.224	1.00	28.81	T6
	ATOM	6213	CB	ASP	A	62	16.370	84.330	16.137	1.00	30.25	T6
	ATOM	6214	CG	ASP	A	62	15.691	83.056	16.572	1.00	25.79	T6
	ATOM	6215	OD1	ASP	A	62	14.445	82.977	16.388	1.00	29.03	T6
30	ATOM	6216	OD2	ASP	A	62	16.400	82.146	17.081	1.00	31.01	T6
	ATOM	6217	C	ASP	A	62	18.569	84.152	17.341	1.00	30.73	T6
	ATOM	6218	O	ASP	A	62	19.532	84.453	16.637	1.00	34.18	T6
	ATOM	6219	N	LYS	A	63	18.625	83.159	18.221	1.00	35.92	T6
	ATOM	6220	CA	LYS	A	63	19.870	82.409	18.368	1.00	31.87	T6
35	ATOM	6221	CB	LYS	A	63	20.286	82.344	19.839	1.00	25.19	T6
	ATOM	6222	CG	LYS	A	63	19.268	81.671	20.741	1.00	35.59	T6
	ATOM	6223	CD	LYS	A	63	19.761	81.610	22.186	1.00	31.92	T6
	ATOM	6224	CE	LYS	A	63	20.013	83.008	22.769	1.00	28.15	T6
	ATOM	6225	NZ	LYS	A	63	20.516	82.980	24.184	1.00	34.70	T6
40	ATOM	6226	C	LYS	A	63	19.830	80.996	17.804	1.00	37.28	T6
	ATOM	6227	O	LYS	A	63	20.649	80.154	18.191	1.00	29.25	T6
	ATOM	6228	N	THR	A	64	18.910	80.735	16.881	1.00	38.61	T6
	ATOM	6229	CA	THR	A	64	18.824	79.394	16.333	1.00	29.51	T6
	ATOM	6230	CB	THR	A	64	17.364	79.043	15.916	1.00	31.96	T6
45	ATOM	6231	OG1	THR	A	64	16.830	80.064	15.065	1.00	35.88	T6
	ATOM	6232	CG2	THR	A	64	16.488	78.899	17.161	1.00	29.87	T6
	ATOM	6233	C	THR	A	64	19.790	79.040	15.192	1.00	29.62	T6
	ATOM	6234	O	THR	A	64	19.372	78.686	14.097	1.00	30.63	T6
	ATOM	6235	N	TYR	A	65	21.085	79.135	15.476	1.00	29.63	T6
50	ATOM	6236	CA	TYR	A	65	22.148	78.781	14.534	1.00	29.62	T6
	ATOM	6237	CB	TYR	A	65	22.334	77.252	14.534	1.00	31.64	T6
	ATOM	6238	CG	TYR	A	65	21.747	76.535	13.338	1.00	35.58	T6
	ATOM	6239	CD1	TYR	A	65	22.508	76.297	12.193	1.00	28.96	T6
	ATOM	6240	CE1	TYR	A	65	21.954	75.646	11.080	1.00	30.06	T6
55	ATOM	6241	CD2	TYR	A	65	20.423	76.109	13.348	1.00	27.54	T6
	ATOM	6242	CE2	TYR	A	65	19.854	75.460	12.244	1.00	30.17	T6
	ATOM	6243	CZ	TYR	A	65	20.619	75.232	11.118	1.00	25.56	T6
	ATOM	6244	OH	TYR	A	65	20.040	74.592	10.040	1.00	32.41	T6
	ATOM	6245	C	TYR	A	65	22.080	79.275	13.081	1.00	34.89	T6
60	ATOM	6246	O	TYR	A	65	23.118	79.371	12.405	1.00	27.83	T6
	ATOM	6247	N	ALA	A	66	20.889	79.574	12.583	1.00	32.75	T6
	ATOM	6248	CA	ALA	A	66	20.783	80.043	11.213	1.00	24.11	T6
	ATOM	6249	CB	ALA	A	66	20.972	78.875	10.254	1.00	25.21	T6
	ATOM	6250	C	ALA	A	66	19.453	80.732	10.954	1.00	29.16	T6
65	ATOM	6251	O	ALA	A	66	18.394	80.128	11.080	1.00	30.71	T6
	ATOM	6252	N	MET	A	67	19.512	82.007	10.598	1.00	31.15	T6

	ATOM	6253	CA	MET	A	67	18.306	82.762	10.301	1.00	26.54	T6
	ATOM	6254	CB	MET	A	67	18.155	83.917	11.285	1.00	32.44	T6
	ATOM	6255	CG	MET	A	67	17.825	83.478	12.704	1.00	37.78	T6
	ATOM	6256	SD	MET	A	67	16.274	82.554	12.782	1.00	37.38	T6
5	ATOM	6257	CE	MET	A	67	15.070	83.896	12.734	1.00	34.97	T6
	ATOM	6258	C	MET	A	67	18.388	83.296	8.880	1.00	26.11	T6
	ATOM	6259	O	MET	A	67	19.441	83.237	8.252	1.00	34.51	T6
	ATOM	6260	N	GLY	A	68	17.272	83.806	8.374	1.00	28.70	T6
	ATOM	6261	CA	GLY	A	68	17.258	84.338	7.025	1.00	28.42	T6
10	ATOM	6262	C	GLY	A	68	15.855	84.619	6.531	1.00	31.99	T6
	ATOM	6263	O	GLY	A	68	14.879	84.237	7.173	1.00	34.25	T6
	ATOM	6264	N	HIS	A	69	15.744	85.301	5.397	1.00	34.06	T6
	ATOM	6265	CA	HIS	A	69	14.435	85.606	4.836	1.00	26.41	T6
	ATOM	6266	CB	HIS	A	69	14.018	87.053	5.148	1.00	34.32	T6
15	ATOM	6267	CG	HIS	A	69	15.017	88.085	4.731	1.00	35.77	T6
	ATOM	6268	CD2	HIS	A	69	15.027	88.939	3.683	1.00	20.19	T6
	ATOM	6269	ND1	HIS	A	69	16.160	88.349	5.452	1.00	26.96	T6
	ATOM	6270	CE1	HIS	A	69	16.830	89.325	4.869	1.00	35.23	T6
	ATOM	6271	NE2	HIS	A	69	16.164	89.701	3.793	1.00	34.06	T6
20	ATOM	6272	C	HIS	A	69	14.361	85.360	3.334	1.00	28.68	T6
	ATOM	6273	O	HIS	A	69	15.381	85.156	2.672	1.00	30.90	T6
	ATOM	6274	N	LEU	A	70	13.138	85.375	2.811	1.00	27.67	T6
	ATOM	6275	CA	LEU	A	70	12.891	85.146	1.397	1.00	30.38	T6
	ATOM	6276	CB	LEU	A	70	12.053	83.885	1.206	1.00	30.53	T6
25	ATOM	6277	CG	LEU	A	70	12.345	82.671	2.078	1.00	27.11	T6
	ATOM	6278	CD1	LEU	A	70	11.269	81.644	1.852	1.00	30.90	T6
	ATOM	6279	CD2	LEU	A	70	13.707	82.102	1.747	1.00	27.43	T6
	ATOM	6280	C	LEU	A	70	12.112	86.297	0.802	1.00	30.14	T6
	ATOM	6281	O	LEU	A	70	11.261	86.884	1.461	1.00	32.02	T6
30	ATOM	6282	N	ILE	A	71	12.409	86.623	-0.445	1.00	41.41	T6
	ATOM	6283	CA	ILE	A	71	11.663	87.659	-1.144	1.00	27.51	T6
	ATOM	6284	CB	ILE	A	71	12.592	88.674	-1.817	1.00	31.89	T6
	ATOM	6285	CG2	ILE	A	71	11.810	89.534	-2.775	1.00	23.45	T6
	ATOM	6286	CG1	ILE	A	71	13.245	89.551	-0.749	1.00	33.44	T6
35	ATOM	6287	CD1	ILE	A	71	14.203	90.571	-1.301	1.00	34.02	T6
	ATOM	6288	C	ILE	A	71	10.885	86.865	-2.185	1.00	32.62	T6
	ATOM	6289	O	ILE	A	71	11.459	86.336	-3.136	1.00	30.88	T6
	ATOM	6290	N	GLN	A	72	9.574	86.763	-2.000	1.00	33.81	T6
	ATOM	6291	CA	GLN	A	72	8.771	85.971	-2.917	1.00	35.24	T6
40	ATOM	6292	CB	GLN	A	72	7.983	84.947	-2.120	1.00	21.95	T6
	ATOM	6293	CG	GLN	A	72	8.835	84.261	-1.081	1.00	28.42	T6
	ATOM	6294	CD	GLN	A	72	8.127	83.111	-0.430	1.00	30.44	T6
	ATOM	6295	OE1	GLN	A	72	7.034	83.269	0.111	1.00	26.04	T6
	ATOM	6296	NE2	GLN	A	72	8.745	81.937	-0.475	1.00	25.95	T6
45	ATOM	6297	C	GLN	A	72	7.834	86.725	-3.843	1.00	33.02	T6
	ATOM	6298	O	GLN	A	72	7.428	87.854	-3.575	1.00	33.13	T6
	ATOM	6299	N	ARG	A	73	7.490	86.062	-4.938	1.00	30.28	T6
	ATOM	6300	CA	ARG	A	73	6.608	86.614	-5.952	1.00	30.16	T6
	ATOM	6301	CB	ARG	A	73	7.375	86.729	-7.264	1.00	39.91	T6
50	ATOM	6302	CG	ARG	A	73	6.562	87.273	-8.388	1.00	31.07	T6
	ATOM	6303	CD	ARG	A	73	7.108	86.828	-9.701	1.00	34.03	T6
	ATOM	6304	NE	ARG	A	73	6.310	87.345	-10.803	1.00	37.02	T6
	ATOM	6305	CZ	ARG	A	73	6.405	86.912	-12.051	1.00	30.54	T6
	ATOM	6306	NH1	ARG	A	73	7.261	85.949	-12.353	1.00	24.06	T6
55	ATOM	6307	NH2	ARG	A	73	5.651	87.448	-12.992	1.00	31.04	T6
	ATOM	6308	C	ARG	A	73	5.374	85.722	-6.164	1.00	28.77	T6
	ATOM	6309	O	ARG	A	73	5.500	84.500	-6.295	1.00	29.80	T6
	ATOM	6310	N	LYS	A	74	4.188	86.334	-6.191	1.00	29.45	T6
	ATOM	6311	CA	LYS	A	74	2.947	85.594	-6.426	1.00	31.49	T6
60	ATOM	6312	CB	LYS	A	74	1.864	86.000	-5.431	1.00	31.47	T6
	ATOM	6313	CG	LYS	A	74	2.138	85.579	-3.995	1.00	35.47	T6
	ATOM	6314	CD	LYS	A	74	1.064	86.106	-3.016	1.00	30.02	T6
	ATOM	6315	CE	LYS	A	74	1.374	85.699	-1.561	1.00	25.45	T6
	ATOM	6316	NZ	LYS	A	74	0.339	86.160	-0.576	1.00	31.26	T6
65	ATOM	6317	C	LYS	A	74	2.465	85.909	-7.828	1.00	33.66	T6
	ATOM	6318	O	LYS	A	74	1.882	86.965	-8.061	1.00	35.00	T6

	ATOM	6319	N	LYS	A	75	2.716	84.995	-8.758	1.00	35.54	T6
	ATOM	6320	CA	LYS	A	75	2.316	85.183	-10.145	1.00	28.66	T6
	ATOM	6321	CB	LYS	A	75	2.747	83.988	-10.990	1.00	32.55	T6
	ATOM	6322	CG	LYS	A	75	4.226	83.712	-11.040	1.00	32.97	T6
5	ATOM	6323	CD	LYS	A	75	4.494	82.492	-11.917	1.00	35.69	T6
	ATOM	6324	CE	LYS	A	75	5.989	82.165	-11.983	1.00	30.48	T6
	ATOM	6325	NZ	LYS	A	75	6.327	80.912	-12.755	1.00	28.30	T6
	ATOM	6326	C	LYS	A	75	0.806	85.339	-10.274	1.00	28.59	T6
	ATOM	6327	O	LYS	A	75	0.052	84.604	-9.644	1.00	35.71	T6
10	ATOM	6328	N	VAL	A	76	0.368	86.290	-11.097	1.00	30.81	T6
	ATOM	6329	CA	VAL	A	76	-1.061	86.507	-11.321	1.00	24.10	T6
	ATOM	6330	CB	VAL	A	76	-1.359	87.837	-12.001	1.00	33.40	T6
	ATOM	6331	CG1	VAL	A	76	-2.805	88.170	-11.830	1.00	29.69	T6
	ATOM	6332	CG2	VAL	A	76	-0.501	88.906	-11.450	1.00	28.57	T6
15	ATOM	6333	C	VAL	A	76	-1.516	85.445	-12.303	1.00	34.50	T6
	ATOM	6334	O	VAL	A	76	-2.599	84.872	-12.172	1.00	28.90	T6
	ATOM	6335	N	HIS	A	77	-0.673	85.212	-13.303	1.00	27.24	T6
	ATOM	6336	CA	HIS	A	77	-0.949	84.233	-14.334	1.00	27.54	T6
	ATOM	6337	CB	HIS	A	77	-0.642	84.834	-15.699	1.00	29.53	T6
20	ATOM	6338	CG	HIS	A	77	-1.414	86.083	-15.989	1.00	28.24	T6
	ATOM	6339	CD2	HIS	A	77	-1.190	87.085	-16.871	1.00	28.43	T6
	ATOM	6340	ND1	HIS	A	77	-2.614	86.371	-15.377	1.00	33.86	T6
	ATOM	6341	CE1	HIS	A	77	-3.102	87.494	-15.872	1.00	31.83	T6
	ATOM	6342	NE2	HIS	A	77	-2.258	87.946	-16.781	1.00	24.75	T6
25	ATOM	6343	C	HIS	A	77	-0.106	82.995	-14.086	1.00	33.68	T6
	ATOM	6344	O	HIS	A	77	1.016	83.099	-13.600	1.00	34.22	T6
	ATOM	6345	N	VAL	A	78	-0.635	81.824	-14.434	1.00	28.05	T6
	ATOM	6346	CA	VAL	A	78	0.097	80.609	-14.166	1.00	32.75	T6
	ATOM	6347	CB	VAL	A	78	-0.585	79.849	-13.015	1.00	29.05	T6
30	ATOM	6348	CG1	VAL	A	78	0.238	78.649	-12.608	1.00	34.85	T6
	ATOM	6349	CG2	VAL	A	78	-0.719	80.774	-11.818	1.00	21.52	T6
	ATOM	6350	C	VAL	A	78	0.447	79.647	-15.307	1.00	33.08	T6
	ATOM	6351	O	VAL	A	78	1.629	79.488	-15.617	1.00	34.78	T6
	ATOM	6352	N	PHE	A	79	-0.522	78.993	-15.935	1.00	26.95	T6
35	ATOM	6353	CA	PHE	A	79	-0.195	78.040	-17.024	1.00	29.55	T6
	ATOM	6354	CB	PHE	A	79	0.872	78.585	-17.990	1.00	22.98	T6
	ATOM	6355	CG	PHE	A	79	0.569	79.940	-18.542	1.00	33.56	T6
	ATOM	6356	CD1	PHE	A	79	1.176	81.074	-18.013	1.00	32.92	T6
	ATOM	6357	CD2	PHE	A	79	-0.327	80.087	-19.587	1.00	34.08	T6
40	ATOM	6358	CE1	PHE	A	79	0.895	82.328	-18.514	1.00	27.74	T6
	ATOM	6359	CE2	PHE	A	79	-0.618	81.342	-20.099	1.00	28.02	T6
	ATOM	6360	CZ	PHE	A	79	-0.006	82.465	-19.561	1.00	27.38	T6
	ATOM	6361	C	PHE	A	79	0.329	76.676	-16.552	1.00	30.49	T6
	ATOM	6362	O	PHE	A	79	1.336	76.587	-15.846	1.00	32.98	T6
45	ATOM	6363	N	GLY	A	80	-0.349	75.616	-16.984	1.00	25.69	T6
	ATOM	6364	CA	GLY	A	80	0.054	74.261	-16.646	1.00	30.52	T6
	ATOM	6365	C	GLY	A	80	0.400	73.983	-15.197	1.00	30.81	T6
	ATOM	6366	O	GLY	A	80	-0.357	74.323	-14.280	1.00	29.73	T6
	ATOM	6367	N	ASP	A	81	1.561	73.361	-14.997	1.00	23.31	T6
50	ATOM	6368	CA	ASP	A	81	2.030	72.995	-13.664	1.00	29.74	T6
	ATOM	6369	CB	ASP	A	81	2.634	71.584	-13.688	1.00	34.29	T6
	ATOM	6370	CG	ASP	A	81	3.895	71.502	-14.517	1.00	36.59	T6
	ATOM	6371	OD1	ASP	A	81	4.144	72.421	-15.320	1.00	25.70	T6
	ATOM	6372	OD2	ASP	A	81	4.638	70.512	-14.380	1.00	27.88	T6
55	ATOM	6373	C	ASP	A	81	3.023	73.975	-13.051	1.00	30.96	T6
	ATOM	6374	O	ASP	A	81	3.881	73.584	-12.259	1.00	29.41	T6
	ATOM	6375	N	GLU	A	82	2.913	75.246	-13.418	1.00	30.44	T6
	ATOM	6376	CA	GLU	A	82	3.793	76.251	-12.844	1.00	33.14	T6
	ATOM	6377	CB	GLU	A	82	3.615	77.601	-13.529	1.00	26.68	T6
60	ATOM	6378	CG	GLU	A	82	4.295	77.792	-14.838	1.00	35.24	T6
	ATOM	6379	CD	GLU	A	82	4.620	79.255	-15.046	1.00	33.59	T6
	ATOM	6380	OE1	GLU	A	82	3.749	80.102	-14.744	1.00	38.30	T6
	ATOM	6381	OE2	GLU	A	82	5.744	79.570	-15.501	1.00	32.48	T6
	ATOM	6382	C	GLU	A	82	3.379	76.448	-11.393	1.00	30.09	T6
65	ATOM	6383	O	GLU	A	82	2.209	76.277	-11.068	1.00	24.25	T6
	ATOM	6384	N	LEU	A	83	4.317	76.796	-10.519	1.00	22.03	T6

	ATOM	6385	CA	LEU	A	83	3.957	77.073	-9.135	1.00	30.15	T6
	ATOM	6386	CB	LEU	A	83	5.091	76.735	-8.171	1.00	31.19	T6
	ATOM	6387	CG	LEU	A	83	5.424	75.261	-7.981	1.00	30.77	T6
	ATOM	6388	CD1	LEU	A	83	5.931	74.680	-9.286	1.00	22.80	T6
5	ATOM	6389	CD2	LEU	A	83	6.474	75.125	-6.914	1.00	31.09	T6
	ATOM	6390	C	LEU	A	83	3.753	78.573	-9.178	1.00	26.36	T6
	ATOM	6391	O	LEU	A	83	4.598	79.294	-9.707	1.00	34.19	T6
	ATOM	6392	N	SER	A	84	2.634	79.048	-8.646	1.00	24.97	T6
	ATOM	6393	CA	SER	A	84	2.352	80.473	-8.679	1.00	27.23	T6
10	ATOM	6394	CB	SER	A	84	0.851	80.719	-8.516	1.00	36.82	T6
	ATOM	6395	OG	SER	A	84	0.331	79.966	-7.442	1.00	29.35	T6
	ATOM	6396	C	SER	A	84	3.130	81.269	-7.652	1.00	35.98	T6
	ATOM	6397	O	SER	A	84	3.059	82.491	-7.636	1.00	28.27	T6
	ATOM	6398	N	LEU	A	85	3.882	80.579	-6.804	1.00	33.63	T6
15	ATOM	6399	CA	LEU	A	85	4.682	81.248	-5.782	1.00	31.95	T6
	ATOM	6400	CB	LEU	A	85	4.280	80.745	-4.393	1.00	30.26	T6
	ATOM	6401	CG	LEU	A	85	4.687	81.534	-3.143	1.00	32.00	T6
	ATOM	6402	CD1	LEU	A	85	6.182	81.442	-2.889	1.00	27.15	T6
	ATOM	6403	CD2	LEU	A	85	4.257	82.981	-3.310	1.00	29.35	T6
20	ATOM	6404	C	LEU	A	85	6.160	80.960	-6.049	1.00	32.51	T6
	ATOM	6405	O	LEU	A	85	6.635	79.847	-5.838	1.00	24.64	T6
	ATOM	6406	N	VAL	A	86	6.881	81.964	-6.537	1.00	30.38	T6
	ATOM	6407	CA	VAL	A	86	8.303	81.811	-6.833	1.00	26.23	T6
	ATOM	6408	CB	VAL	A	86	8.666	82.360	-8.225	1.00	34.51	T6
25	ATOM	6409	CG1	VAL	A	86	10.167	82.281	-8.428	1.00	23.74	T6
	ATOM	6410	CG2	VAL	A	86	7.944	81.583	-9.306	1.00	38.73	T6
	ATOM	6411	C	VAL	A	86	9.089	82.607	-5.822	1.00	25.67	T6
	ATOM	6412	O	VAL	A	86	8.710	83.720	-5.476	1.00	29.66	T6
	ATOM	6413	N	THR	A	87	10.180	82.041	-5.333	1.00	33.00	T6
30	ATOM	6414	CA	THR	A	87	10.981	82.771	-4.380	1.00	36.13	T6
	ATOM	6415	CB	THR	A	87	11.287	81.927	-3.129	1.00	25.17	T6
	ATOM	6416	OG1	THR	A	87	12.692	81.720	-3.023	1.00	35.41	T6
	ATOM	6417	CG2	THR	A	87	10.569	80.589	-3.200	1.00	34.69	T6
	ATOM	6418	C	THR	A	87	12.258	83.217	-5.075	1.00	28.75	T6
35	ATOM	6419	O	THR	A	87	13.029	82.408	-5.572	1.00	40.47	T6
	ATOM	6420	N	LEU	A	88	12.439	84.529	-5.145	1.00	25.43	T6
	ATOM	6421	CA	LEU	A	88	13.602	85.146	-5.767	1.00	34.43	T6
	ATOM	6422	CB	LEU	A	88	13.187	86.459	-6.434	1.00	33.36	T6
	ATOM	6423	CG	LEU	A	88	12.089	86.602	-7.498	1.00	27.30	T6
40	ATOM	6424	CD1	LEU	A	88	11.044	85.556	-7.345	1.00	31.67	T6
	ATOM	6425	CD2	LEU	A	88	11.463	87.965	-7.369	1.00	27.20	T6
	ATOM	6426	C	LEU	A	88	14.579	85.488	-4.647	1.00	24.15	T6
	ATOM	6427	O	LEU	A	88	14.174	85.818	-3.525	1.00	26.55	T6
	ATOM	6428	N	PHE	A	89	15.868	85.398	-4.900	1.00	31.31	T6
45	ATOM	6429	CA	PHE	A	89	16.822	85.817	-3.859	1.00	32.54	T6
	ATOM	6430	CB	PHE	A	89	16.555	87.292	-3.525	1.00	37.42	T6
	ATOM	6431	CG	PHE	A	89	16.291	88.123	-4.745	1.00	30.41	T6
	ATOM	6432	CD1	PHE	A	89	15.388	89.177	-4.708	1.00	34.53	T6
	ATOM	6433	CD2	PHE	A	89	16.882	87.774	-5.980	1.00	35.96	T6
50	ATOM	6434	CE1	PHE	A	89	15.062	89.867	-5.886	1.00	27.75	T6
	ATOM	6435	CE2	PHE	A	89	16.568	88.452	-7.162	1.00	27.18	T6
	ATOM	6436	CZ	PHE	A	89	15.654	89.499	-7.121	1.00	29.26	T6
	ATOM	6437	C	PHE	A	89	16.951	85.005	-2.568	1.00	28.38	T6
	ATOM	6438	O	PHE	A	89	17.555	83.934	-2.588	1.00	32.73	T6
55	ATOM	6439	N	ARG	A	90	16.435	85.499	-1.443	1.00	29.48	T6
	ATOM	6440	CA	ARG	A	90	16.601	84.747	-0.186	1.00	29.78	T6
	ATOM	6441	CB	ARG	A	90	16.096	83.315	-0.362	1.00	28.58	T6
	ATOM	6442	CG	ARG	A	90	16.765	82.294	0.547	1.00	26.96	T6
	ATOM	6443	CD	ARG	A	90	16.522	80.885	0.048	1.00	27.92	T6
60	ATOM	6444	NE	ARG	A	90	17.345	79.933	0.778	1.00	35.05	T6
	ATOM	6445	CZ	ARG	A	90	17.906	78.853	0.239	1.00	29.91	T6
	ATOM	6446	NH1	ARG	A	90	17.737	78.568	-1.048	1.00	34.52	T6
	ATOM	6447	NH2	ARG	A	90	18.663	78.064	0.992	1.00	36.44	T6
	ATOM	6448	C	ARG	A	90	18.059	84.674	0.340	1.00	29.81	T6
65	ATOM	6449	O	ARG	A	90	18.973	84.278	-0.382	1.00	26.68	T6
	ATOM	6450	N	CYS	A	91	18.257	85.019	1.611	1.00	22.96	T6

	ATOM	6451	CA	CYS	A	91	19.582	84.988	2.229	1.00	34.06	T6
	ATOM	6452	CB	CYS	A	91	20.078	86.414	2.441	1.00	31.56	T6
	ATOM	6453	SG	CYS	A	91	18.923	87.414	3.403	1.00	32.97	T6
	ATOM	6454	C	CYS	A	91	19.583	84.229	3.564	1.00	33.08	T6
5	ATOM	6455	O	CYS	A	91	18.526	83.951	4.130	1.00	28.98	T6
	ATOM	6456	N	ILE	A	92	20.775	83.901	4.062	1.00	29.10	T6
	ATOM	6457	CA	ILE	A	92	20.934	83.149	5.313	1.00	32.77	T6
	ATOM	6458	CB	ILE	A	92	21.264	81.681	5.032	1.00	32.13	T6
	ATOM	6459	CG2	ILE	A	92	21.349	80.909	6.330	1.00	35.19	T6
10	ATOM	6460	CG1	ILE	A	92	20.186	81.070	4.138	1.00	31.03	T6
	ATOM	6461	CD1	ILE	A	92	20.575	79.722	3.546	1.00	30.97	T6
	ATOM	6462	C	ILE	A	92	22.086	83.703	6.134	1.00	30.98	T6
	ATOM	6463	O	ILE	A	92	23.010	84.280	5.585	1.00	25.56	T6
	ATOM	6464	N	GLN	A	93	22.038	83.520	7.450	1.00	38.13	T6
15	ATOM	6465	CA	GLN	A	93	23.103	83.997	8.333	1.00	30.94	T6
	ATOM	6466	CB	GLN	A	93	22.857	85.449	8.754	1.00	30.39	T6
	ATOM	6467	CG	GLN	A	93	23.385	86.492	7.779	1.00	27.52	T6
	ATOM	6468	CD	GLN	A	93	24.744	87.040	8.179	1.00	29.34	T6
	ATOM	6469	OE1	GLN	A	93	25.774	86.378	8.019	1.00	30.55	T6
20	ATOM	6470	NE2	GLN	A	93	24.751	88.259	8.712	1.00	34.55	T6
	ATOM	6471	C	GLN	A	93	23.213	83.131	9.577	1.00	35.37	T6
	ATOM	6472	O	GLN	A	93	22.250	82.988	10.335	1.00	30.70	T6
	ATOM	6473	N	ASN	A	94	24.385	82.539	9.780	1.00	33.32	T6
	ATOM	6474	CA	ASN	A	94	24.589	81.710	10.955	1.00	32.00	T6
25	ATOM	6475	CB	ASN	A	94	25.993	81.095	10.949	1.00	21.25	T6
	ATOM	6476	CG	ASN	A	94	26.086	79.863	10.054	1.00	30.31	T6
	ATOM	6477	OD1	ASN	A	94	25.316	78.915	10.215	1.00	29.63	T6
	ATOM	6478	ND2	ASN	A	94	27.029	79.868	9.111	1.00	32.95	T6
	ATOM	6479	C	ASN	A	94	24.421	82.632	12.143	1.00	30.97	T6
30	ATOM	6480	O	ASN	A	94	24.753	83.812	12.060	1.00	30.08	T6
	ATOM	6481	N	MET	A	95	23.884	82.107	13.237	1.00	32.13	T6
	ATOM	6482	CA	MET	A	95	23.683	82.900	14.442	1.00	32.20	T6
	ATOM	6483	CB	MET	A	95	22.218	82.849	14.884	1.00	32.54	T6
	ATOM	6484	CG	MET	A	95	21.244	83.412	13.876	1.00	38.53	T6
35	ATOM	6485	SD	MET	A	95	21.690	85.077	13.308	1.00	27.66	T6
	ATOM	6486	CE	MET	A	95	21.253	86.079	14.718	1.00	27.46	T6
	ATOM	6487	C	MET	A	95	24.554	82.389	15.584	1.00	32.86	T6
	ATOM	6488	O	MET	A	95	24.877	81.202	15.643	1.00	28.00	T6
	ATOM	6489	N	PRO	A	96	24.955	83.288	16.503	1.00	26.14	T6
40	ATOM	6490	CD	PRO	A	96	24.750	84.741	16.465	1.00	23.46	T6
	ATOM	6491	CA	PRO	A	96	25.784	82.929	17.655	1.00	40.53	T6
	ATOM	6492	CB	PRO	A	96	26.278	84.273	18.180	1.00	33.19	T6
	ATOM	6493	CG	PRO	A	96	26.038	85.227	17.054	1.00	28.71	T6
	ATOM	6494	C	PRO	A	96	24.834	82.292	18.647	1.00	26.87	T6
45	ATOM	6495	O	PRO	A	96	23.667	82.020	18.322	1.00	31.48	T6
	ATOM	6496	N	GLU	A	97	25.310	82.079	19.865	1.00	24.68	T6
	ATOM	6497	CA	GLU	A	97	24.464	81.473	20.871	1.00	30.72	T6
	ATOM	6498	CB	GLU	A	97	25.146	80.234	21.415	1.00	26.67	T6
	ATOM	6499	CG	GLU	A	97	24.209	79.341	22.174	1.00	27.69	T6
50	ATOM	6500	CD	GLU	A	97	24.425	77.884	21.812	1.00	36.96	T6
	ATOM	6501	OE1	GLU	A	97	25.563	77.385	22.042	1.00	26.61	T6
	ATOM	6502	OE2	GLU	A	97	23.466	77.245	21.291	1.00	31.79	T6
	ATOM	6503	C	GLU	A	97	24.199	82.459	21.991	1.00	28.15	T6
	ATOM	6504	O	GLU	A	97	23.209	82.359	22.710	1.00	33.87	T6
55	ATOM	6505	N	THR	A	98	25.084	83.436	22.108	1.00	29.96	T6
	ATOM	6506	CA	THR	A	98	24.976	84.420	23.159	1.00	27.97	T6
	ATOM	6507	CB	THR	A	98	26.355	84.810	23.656	1.00	35.19	T6
	ATOM	6508	OG1	THR	A	98	27.077	85.435	22.581	1.00	30.39	T6
	ATOM	6509	CG2	THR	A	98	27.106	83.568	24.141	1.00	29.29	T6
60	ATOM	6510	C	THR	A	98	24.229	85.699	22.804	1.00	32.34	T6
	ATOM	6511	O	THR	A	98	23.104	85.907	23.266	1.00	28.36	T6
	ATOM	6512	N	LEU	A	99	24.826	86.561	21.990	1.00	27.66	T6
	ATOM	6513	CA	LEU	A	99	24.155	87.816	21.697	1.00	28.72	T6
	ATOM	6514	CB	LEU	A	99	25.026	88.972	22.185	1.00	29.99	T6
65	ATOM	6515	CG	LEU	A	99	25.236	88.948	23.700	1.00	38.54	T6
	ATOM	6516	CD1	LEU	A	99	26.221	90.016	24.129	1.00	31.84	T6

	ATOM	6517	CD2	LEU	A	99	23.901	89.161	24.376	1.00	36.87	T6
	ATOM	6518	C	LEU	A	99	23.782	88.009	20.246	1.00	30.77	T6
	ATOM	6519	O	LEU	A	99	24.382	88.832	19.542	1.00	30.33	T6
	ATOM	6520	N	PRO	A	100	22.760	87.270	19.781	1.00	32.52	T6
5	ATOM	6521	CD	PRO	A	100	21.926	86.353	20.577	1.00	31.71	T6
	ATOM	6522	CA	PRO	A	100	22.274	87.332	18.397	1.00	28.07	T6
	ATOM	6523	CB	PRO	A	100	20.946	86.584	18.469	1.00	30.17	T6
	ATOM	6524	CG	PRO	A	100	21.228	85.539	19.497	1.00	36.77	T6
	ATOM	6525	C	PRO	A	100	22.116	88.751	17.878	1.00	25.34	T6
10	ATOM	6526	O	PRO	A	100	21.399	89.563	18.467	1.00	31.62	T6
	ATOM	6527	N	ASN	A	101	22.797	89.036	16.775	1.00	29.03	T6
	ATOM	6528	CA	ASN	A	101	22.751	90.352	16.151	1.00	31.84	T6
	ATOM	6529	CB	ASN	A	101	23.583	91.346	16.939	1.00	28.34	T6
	ATOM	6530	CG	ASN	A	101	22.806	91.988	18.030	1.00	33.01	T6
15	ATOM	6531	OD1	ASN	A	101	21.900	92.775	17.762	1.00	30.84	T6
	ATOM	6532	ND2	ASN	A	101	23.135	91.654	19.281	1.00	36.28	T6
	ATOM	6533	C	ASN	A	101	23.322	90.276	14.763	1.00	29.35	T6
	ATOM	6534	O	ASN	A	101	24.498	90.609	14.563	1.00	27.69	T6
	ATOM	6535	N	ASN	A	102	22.523	89.859	13.788	1.00	35.91	T6
20	ATOM	6536	CA	ASN	A	102	23.110	89.786	12.476	1.00	26.77	T6
	ATOM	6537	CB	ASN	A	102	23.232	88.327	12.032	1.00	27.45	T6
	ATOM	6538	CG	ASN	A	102	24.495	87.678	12.580	1.00	27.75	T6
	ATOM	6539	OD1	ASN	A	102	25.582	88.269	12.530	1.00	32.66	T6
	ATOM	6540	ND2	ASN	A	102	24.360	86.468	13.110	1.00	28.19	T6
25	ATOM	6541	C	ASN	A	102	22.630	90.665	11.335	1.00	30.73	T6
	ATOM	6542	O	ASN	A	102	23.475	91.257	10.663	1.00	32.11	T6
	ATOM	6543	N	SER	A	103	21.337	90.799	11.076	1.00	36.07	T6
	ATOM	6544	CA	SER	A	103	20.980	91.672	9.949	1.00	31.41	T6
	ATOM	6545	CB	SER	A	103	21.470	93.110	10.215	1.00	33.29	T6
30	ATOM	6546	OG	SER	A	103	22.147	93.661	9.097	1.00	29.73	T6
	ATOM	6547	C	SER	A	103	21.575	91.159	8.619	1.00	27.59	T6
	ATOM	6548	O	SER	A	103	22.769	90.877	8.497	1.00	31.08	T6
	ATOM	6549	N	CYS	A	104	20.734	91.052	7.607	1.00	28.06	T6
	ATOM	6550	CA	CYS	A	104	21.208	90.558	6.341	1.00	30.10	T6
35	ATOM	6551	CB	CYS	A	104	21.011	89.046	6.302	1.00	29.90	T6
	ATOM	6552	SG	CYS	A	104	21.823	88.216	4.951	1.00	28.44	T6
	ATOM	6553	C	CYS	A	104	20.448	91.236	5.220	1.00	31.95	T6
	ATOM	6554	O	CYS	A	104	19.224	91.259	5.216	1.00	30.79	T6
	ATOM	6555	N	TYR	A	105	21.192	91.797	4.277	1.00	36.50	T6
40	ATOM	6556	CA	TYR	A	105	20.616	92.482	3.129	1.00	37.13	T6
	ATOM	6557	CB	TYR	A	105	21.314	93.829	2.925	1.00	35.26	T6
	ATOM	6558	CG	TYR	A	105	20.868	94.594	1.697	1.00	33.37	T6
	ATOM	6559	CD1	TYR	A	105	19.876	95.565	1.778	1.00	31.73	T6
	ATOM	6560	CE1	TYR	A	105	19.482	96.288	0.658	1.00	27.23	T6
45	ATOM	6561	CD2	TYR	A	105	21.453	94.359	0.455	1.00	26.51	T6
	ATOM	6562	CE2	TYR	A	105	21.061	95.075	-0.675	1.00	32.65	T6
	ATOM	6563	CZ	TYR	A	105	20.080	96.038	-0.562	1.00	27.12	T6
	ATOM	6564	OH	TYR	A	105	19.716	96.769	-1.663	1.00	28.23	T6
	ATOM	6565	C	TYR	A	105	20.801	91.635	1.882	1.00	30.28	T6
50	ATOM	6566	O	TYR	A	105	21.794	90.932	1.744	1.00	41.27	T6
	ATOM	6567	N	SER	A	106	19.840	91.703	0.974	1.00	34.24	T6
	ATOM	6568	CA	SER	A	106	19.930	90.965	-0.273	1.00	27.57	T6
	ATOM	6569	CB	SER	A	106	19.541	89.497	-0.065	1.00	23.29	T6
	ATOM	6570	OG	SER	A	106	19.865	88.720	-1.205	1.00	23.89	T6
55	ATOM	6571	C	SER	A	106	18.987	91.640	-1.253	1.00	31.80	T6
	ATOM	6572	O	SER	A	106	17.930	92.128	-0.859	1.00	29.16	T6
	ATOM	6573	N	ALA	A	107	19.380	91.681	-2.523	1.00	27.45	T6
	ATOM	6574	CA	ALA	A	107	18.572	92.316	-3.550	1.00	29.47	T6
	ATOM	6575	CB	ALA	A	107	18.819	93.807	-3.542	1.00	27.95	T6
60	ATOM	6576	C	ALA	A	107	18.884	91.748	-4.922	1.00	33.57	T6
	ATOM	6577	O	ALA	A	107	19.898	91.085	-5.110	1.00	34.73	T6
	ATOM	6578	N	GLY	A	108	17.998	92.014	-5.876	1.00	28.30	T6
	ATOM	6579	CA	GLY	A	108	18.190	91.529	-7.231	1.00	26.99	T6
	ATOM	6580	C	GLY	A	108	17.165	92.117	-8.180	1.00	34.09	T6
65	ATOM	6581	O	GLY	A	108	16.313	92.892	-7.767	1.00	32.96	T6
	ATOM	6582	N	ILE	A	109	17.248	91.757	-9.453	1.00	31.96	T6

	ATOM	6583	CA	ILE	A	109	16.315	92.253	-10.454	1.00	30.94	T6
	ATOM	6584	CB	ILE	A	109	17.059	92.825	-11.665	1.00	40.15	T6
	ATOM	6585	CG2	ILE	A	109	16.071	93.298	-12.706	1.00	30.60	T6
	ATOM	6586	CG1	ILE	A	109	17.952	93.974	-11.224	1.00	31.32	T6
5	ATOM	6587	CD1	ILE	A	109	18.875	94.461	-12.303	1.00	25.03	T6
	ATOM	6588	C	ILE	A	109	15.433	91.106	-10.936	1.00	26.41	T6
	ATOM	6589	O	ILE	A	109	15.902	89.982	-11.100	1.00	31.91	T6
	ATOM	6590	N	ALA	A	110	14.157	91.391	-11.163	1.00	29.32	T6
	ATOM	6591	CA	ALA	A	110	13.224	90.378	-11.631	1.00	31.58	T6
10	ATOM	6592	CB	ALA	A	110	12.572	89.701	-10.460	1.00	33.78	T6
	ATOM	6593	C	ALA	A	110	12.171	91.030	-12.497	1.00	36.12	T6
	ATOM	6594	O	ALA	A	110	11.913	92.223	-12.367	1.00	30.44	T6
	ATOM	6595	N	LYS	A	111	11.576	90.260	-13.401	1.00	29.94	T6
	ATOM	6596	CA	LYS	A	111	10.533	90.812	-14.243	1.00	28.87	T6
15	ATOM	6597	CB	LYS	A	111	10.573	90.215	-15.636	1.00	33.94	T6
	ATOM	6598	CG	LYS	A	111	9.515	90.832	-16.544	1.00	42.23	T6
	ATOM	6599	CD	LYS	A	111	9.592	90.288	-17.968	1.00	29.56	T6
	ATOM	6600	CE	LYS	A	111	8.540	90.933	-18.874	1.00	29.14	T6
	ATOM	6601	NZ	LYS	A	111	8.615	90.403	-20.283	1.00	26.87	T6
20	ATOM	6602	C	LYS	A	111	9.207	90.479	-13.587	1.00	25.97	T6
	ATOM	6603	O	LYS	A	111	8.982	89.332	-13.204	1.00	22.43	T6
	ATOM	6604	N	LEU	A	112	8.340	91.481	-13.456	1.00	31.81	T6
	ATOM	6605	CA	LEU	A	112	7.036	91.303	-12.821	1.00	28.82	T6
	ATOM	6606	CB	LEU	A	112	7.003	92.077	-11.507	1.00	29.38	T6
25	ATOM	6607	CG	LEU	A	112	8.177	91.872	-10.558	1.00	29.99	T6
	ATOM	6608	CD1	LEU	A	112	8.148	92.930	-9.469	1.00	33.07	T6
	ATOM	6609	CD2	LEU	A	112	8.106	90.479	-9.972	1.00	29.29	T6
	ATOM	6610	C	LEU	A	112	5.908	91.814	-13.723	1.00	24.60	T6
	ATOM	6611	O	LEU	A	112	6.154	92.595	-14.645	1.00	31.42	T6
30	ATOM	6612	N	GLU	A	113	4.675	91.382	-13.441	1.00	27.19	T6
	ATOM	6613	CA	GLU	A	113	3.511	91.803	-14.218	1.00	32.40	T6
	ATOM	6614	CB	GLU	A	113	2.911	90.642	-14.986	1.00	37.44	T6
	ATOM	6615	CG	GLU	A	113	3.875	89.689	-15.597	1.00	39.66	T6
	ATOM	6616	CD	GLU	A	113	3.218	88.887	-16.705	1.00	28.86	T6
35	ATOM	6617	OE1	GLU	A	113	2.062	88.409	-16.494	1.00	32.31	T6
	ATOM	6618	OE2	GLU	A	113	3.860	88.741	-17.788	1.00	33.63	T6
	ATOM	6619	C	GLU	A	113	2.397	92.352	-13.351	1.00	27.45	T6
	ATOM	6620	O	GLU	A	113	2.312	92.041	-12.158	1.00	33.12	T6
	ATOM	6621	N	GLU	A	114	1.525	93.145	-13.968	1.00	32.92	T6
40	ATOM	6622	CA	GLU	A	114	0.373	93.713	-13.277	1.00	26.39	T6
	ATOM	6623	CB	GLU	A	114	-0.667	94.188	-14.265	1.00	29.18	T6
	ATOM	6624	CG	GLU	A	114	-0.538	95.589	-14.740	1.00	36.82	T6
	ATOM	6625	CD	GLU	A	114	-1.854	96.069	-15.322	1.00	36.60	T6
	ATOM	6626	OE1	GLU	A	114	-2.844	96.150	-14.551	1.00	37.24	T6
45	ATOM	6627	OE2	GLU	A	114	-1.907	96.347	-16.546	1.00	32.38	T6
	ATOM	6628	C	GLU	A	114	-0.303	92.643	-12.447	1.00	26.23	T6
	ATOM	6629	O	GLU	A	114	-0.691	91.607	-12.977	1.00	27.11	T6
	ATOM	6630	N	GLY	A	115	-0.465	92.890	-11.157	1.00	32.29	T6
	ATOM	6631	CA	GLY	A	115	-1.137	91.912	-10.331	1.00	24.29	T6
50	ATOM	6632	C	GLY	A	115	-0.212	91.048	-9.507	1.00	26.50	T6
	ATOM	6633	O	GLY	A	115	-0.658	90.403	-8.552	1.00	32.46	T6
	ATOM	6634	N	ASP	A	116	1.065	90.992	-9.871	1.00	27.99	T6
	ATOM	6635	CA	ASP	A	116	1.983	90.184	-9.085	1.00	33.46	T6
	ATOM	6636	CB	ASP	A	116	3.364	90.109	-9.742	1.00	36.48	T6
55	ATOM	6637	CG	ASP	A	116	3.380	89.257	-11.003	1.00	29.09	T6
	ATOM	6638	OD1	ASP	A	116	2.536	88.346	-11.124	1.00	31.26	T6
	ATOM	6639	OD2	ASP	A	116	4.262	89.486	-11.865	1.00	25.86	T6
	ATOM	6640	C	ASP	A	116	2.114	90.848	-7.719	1.00	30.22	T6
	ATOM	6641	O	ASP	A	116	1.924	92.058	-7.586	1.00	32.03	T6
60	ATOM	6642	N	GLU	A	117	2.424	90.057	-6.701	1.00	32.55	T6
	ATOM	6643	CA	GLU	A	117	2.601	90.608	-5.372	1.00	30.53	T6
	ATOM	6644	CB	GLU	A	117	1.503	90.131	-4.436	1.00	30.95	T6
	ATOM	6645	CG	GLU	A	117	0.111	90.409	-4.935	1.00	30.92	T6
	ATOM	6646	CD	GLU	A	117	-0.950	89.997	-3.930	1.00	33.53	T6
65	ATOM	6647	OE1	GLU	A	117	-0.759	88.954	-3.252	1.00	39.54	T6
	ATOM	6648	OE2	GLU	A	117	-1.975	90.712	-3.829	1.00	27.83	T6

	ATOM	6649	C	GLU A 117	3.942	90.146	-4.843	1.00	24.51	T6
	ATOM	6650	O	GLU A 117	4.348	89.003	-5.074	1.00	32.37	T6
	ATOM	6651	N	LEU A 118	4.633	91.045	-4.148	1.00	30.53	T6
	ATOM	6652	CA	LEU A 118	5.925	90.730	-3.562	1.00	34.16	T6
5	ATOM	6653	CB	LEU A 118	6.948	91.809	-3.900	1.00	35.66	T6
	ATOM	6654	CG	LEU A 118	7.335	91.965	-5.366	1.00	35.30	T6
	ATOM	6655	CD1	LEU A 118	8.411	93.016	-5.492	1.00	36.79	T6
	ATOM	6656	CD2	LEU A 118	7.831	90.642	-5.904	1.00	30.30	T6
	ATOM	6657	C	LEU A 118	5.758	90.665	-2.059	1.00	23.81	T6
10	ATOM	6658	O	LEU A 118	4.964	91.407	-1.493	1.00	26.11	T6
	ATOM	6659	N	GLN A 119	6.498	89.776	-1.413	1.00	31.55	T6
	ATOM	6660	CA	GLN A 119	6.435	89.658	0.034	1.00	34.30	T6
	ATOM	6661	CB	GLN A 119	5.310	88.718	0.425	1.00	23.78	T6
	ATOM	6662	CG	GLN A 119	5.505	87.306	-0.071	1.00	31.21	T6
15	ATOM	6663	CD	GLN A 119	4.326	86.407	0.264	1.00	27.43	T6
	ATOM	6664	OE1	GLN A 119	4.442	85.178	0.238	1.00	36.39	T6
	ATOM	6665	NE2	GLN A 119	3.177	87.016	0.567	1.00	26.42	T6
	ATOM	6666	C	GLN A 119	7.764	89.172	0.617	1.00	27.27	T6
	ATOM	6667	O	GLN A 119	8.555	88.520	-0.066	1.00	28.83	T6
20	ATOM	6668	N	LEU A 120	7.997	89.509	1.881	1.00	36.29	T6
	ATOM	6669	CA	LEU A 120	9.218	89.148	2.597	1.00	32.92	T6
	ATOM	6670	CB	LEU A 120	9.785	90.403	3.274	1.00	36.00	T6
	ATOM	6671	CG	LEU A 120	11.181	90.509	3.897	1.00	28.81	T6
	ATOM	6672	CD1	LEU A 120	11.476	89.304	4.764	1.00	30.98	T6
25	ATOM	6673	CD2	LEU A 120	12.203	90.639	2.797	1.00	32.96	T6
	ATOM	6674	C	LEU A 120	8.823	88.120	3.655	1.00	28.67	T6
	ATOM	6675	O	LEU A 120	8.000	88.409	4.525	1.00	37.67	T6
	ATOM	6676	N	ALA A 121	9.408	86.928	3.602	1.00	37.69	T6
	ATOM	6677	CA	ALA A 121	9.047	85.897	4.569	1.00	35.06	T6
30	ATOM	6678	CB	ALA A 121	8.223	84.822	3.877	1.00	31.80	T6
	ATOM	6679	C	ALA A 121	10.207	85.246	5.311	1.00	26.48	T6
	ATOM	6680	O	ALA A 121	11.268	84.984	4.739	1.00	30.81	T6
	ATOM	6681	N	ILE A 122	9.991	84.981	6.596	1.00	34.87	T6
	ATOM	6682	CA	ILE A 122	10.994	84.319	7.420	1.00	32.00	T6
35	ATOM	6683	CB	ILE A 122	11.197	85.045	8.749	1.00	34.75	T6
	ATOM	6684	CG2	ILE A 122	12.271	84.340	9.554	1.00	32.79	T6
	ATOM	6685	CG1	ILE A 122	11.582	86.503	8.484	1.00	30.41	T6
	ATOM	6686	CD1	ILE A 122	11.809	87.320	9.735	1.00	33.95	T6
	ATOM	6687	C	ILE A 122	10.523	82.889	7.701	1.00	27.34	T6
40	ATOM	6688	O	ILE A 122	9.473	82.668	8.316	1.00	33.89	T6
	ATOM	6689	N	PRO A 123	11.297	81.895	7.239	1.00	24.22	T6
	ATOM	6690	CD	PRO A 123	12.495	82.067	6.409	1.00	27.27	T6
	ATOM	6691	CA	PRO A 123	10.995	80.470	7.411	1.00	37.40	T6
	ATOM	6692	CB	PRO A 123	11.979	79.786	6.466	1.00	34.37	T6
45	ATOM	6693	CG	PRO A 123	12.395	80.873	5.521	1.00	29.99	T6
	ATOM	6694	C	PRO A 123	11.170	79.973	8.843	1.00	26.29	T6
	ATOM	6695	O	PRO A 123	11.917	79.026	9.081	1.00	30.51	T6
	ATOM	6696	N	ARG A 124	10.504	80.622	9.792	1.00	31.30	T6
	ATOM	6697	CA	ARG A 124	10.573	80.213	11.189	1.00	28.18	T6
50	ATOM	6698	CB	ARG A 124	11.677	80.932	11.921	1.00	28.87	T6
	ATOM	6699	CG	ARG A 124	13.008	80.246	11.787	1.00	35.18	T6
	ATOM	6700	CD	ARG A 124	13.693	80.227	13.140	1.00	32.01	T6
	ATOM	6701	NE	ARG A 124	13.076	79.270	14.065	1.00	28.80	T6
	ATOM	6702	CZ	ARG A 124	13.376	79.199	15.358	1.00	24.67	T6
55	ATOM	6703	NH1	ARG A 124	14.273	80.030	15.868	1.00	33.08	T6
	ATOM	6704	NH2	ARG A 124	12.798	78.295	16.134	1.00	33.57	T6
	ATOM	6705	C	ARG A 124	9.260	80.514	11.855	1.00	36.69	T6
	ATOM	6706	O	ARG A 124	8.518	81.383	11.386	1.00	31.90	T6
	ATOM	6707	N	GLU A 125	8.972	79.812	12.948	1.00	31.69	T6
60	ATOM	6708	CA	GLU A 125	7.701	80.012	13.618	1.00	28.96	T6
	ATOM	6709	CB	GLU A 125	7.389	78.827	14.521	1.00	32.39	T6
	ATOM	6710	CG	GLU A 125	7.002	77.582	13.724	1.00	32.65	T6
	ATOM	6711	CD	GLU A 125	5.839	76.817	14.359	1.00	28.14	T6
	ATOM	6712	OE1	GLU A 125	5.952	76.437	15.562	1.00	28.07	T6
65	ATOM	6713	OE2	GLU A 125	4.817	76.600	13.646	1.00	33.19	T6
	ATOM	6714	C	GLU A 125	7.571	81.320	14.373	1.00	26.77	T6

	ATOM	6715	O	GLU	A	125	6.580	82.037	14.200	1.00	33.00	T6
	ATOM	6716	N	ASN	A	126	8.538	81.647	15.215	1.00	28.19	T6
	ATOM	6717	CA	ASN	A	126	8.445	82.917	15.921	1.00	31.05	T6
	ATOM	6718	CB	ASN	A	126	7.947	82.735	17.356	1.00	25.48	T6
5	ATOM	6719	CG	ASN	A	126	6.430	82.608	17.434	1.00	33.41	T6
	ATOM	6720	OD1	ASN	A	126	5.861	81.547	17.172	1.00	27.25	T6
	ATOM	6721	ND2	ASN	A	126	5.766	83.705	17.785	1.00	35.86	T6
	ATOM	6722	C	ASN	A	126	9.801	83.560	15.913	1.00	33.69	T6
	ATOM	6723	O	ASN	A	126	10.415	83.776	16.958	1.00	29.95	T6
10	ATOM	6724	N	ALA	A	127	10.262	83.859	14.707	1.00	29.24	T6
	ATOM	6725	CA	ALA	A	127	11.565	84.462	14.509	1.00	34.14	T6
	ATOM	6726	CB	ALA	A	127	11.692	84.926	13.072	1.00	27.00	T6
	ATOM	6727	C	ALA	A	127	11.810	85.625	15.455	1.00	28.20	T6
	ATOM	6728	O	ALA	A	127	10.949	86.486	15.614	1.00	32.81	T6
15	ATOM	6729	N	GLN	A	128	12.972	85.628	16.102	1.00	33.89	T6
	ATOM	6730	CA	GLN	A	128	13.340	86.718	16.995	1.00	35.23	T6
	ATOM	6731	CB	GLN	A	128	14.349	86.233	18.026	1.00	32.39	T6
	ATOM	6732	CG	GLN	A	128	13.756	85.221	18.987	1.00	32.65	T6
	ATOM	6733	CD	GLN	A	128	12.398	85.669	19.513	1.00	27.07	T6
20	ATOM	6734	OE1	GLN	A	128	12.267	86.735	20.126	1.00	23.24	T6
	ATOM	6735	NE2	GLN	A	128	11.376	84.857	19.264	1.00	28.47	T6
	ATOM	6736	C	GLN	A	128	13.946	87.787	16.096	1.00	25.97	T6
	ATOM	6737	O	GLN	A	128	15.107	87.709	15.698	1.00	30.73	T6
	ATOM	6738	N	ILE	A	129	13.136	88.790	15.788	1.00	34.38	T6
25	ATOM	6739	CA	ILE	A	129	13.510	89.857	14.872	1.00	28.70	T6
	ATOM	6740	CB	ILE	A	129	12.548	89.764	13.633	1.00	29.09	T6
	ATOM	6741	CG2	ILE	A	129	11.896	91.090	13.311	1.00	30.75	T6
	ATOM	6742	CG1	ILE	A	129	13.298	89.174	12.454	1.00	32.56	T6
	ATOM	6743	CD1	ILE	A	129	13.809	87.789	12.736	1.00	31.81	T6
30	ATOM	6744	C	ILE	A	129	13.470	91.254	15.493	1.00	31.86	T6
	ATOM	6745	O	ILE	A	129	12.827	91.465	16.516	1.00	29.31	T6
	ATOM	6746	N	SER	A	130	14.174	92.201	14.887	1.00	33.70	T6
	ATOM	6747	CA	SER	A	130	14.141	93.575	15.365	1.00	32.85	T6
	ATOM	6748	CB	SER	A	130	15.525	94.208	15.320	1.00	24.82	T6
35	ATOM	6749	OG	SER	A	130	15.435	95.612	15.531	1.00	33.44	T6
	ATOM	6750	C	SER	A	130	13.221	94.343	14.426	1.00	28.25	T6
	ATOM	6751	O	SER	A	130	13.412	94.315	13.213	1.00	33.11	T6
	ATOM	6752	N	LEU	A	131	12.224	95.031	14.969	1.00	28.62	T6
	ATOM	6753	CA	LEU	A	131	11.298	95.778	14.120	1.00	28.62	T6
40	ATOM	6754	CB	LEU	A	131	9.859	95.597	14.607	1.00	25.60	T6
	ATOM	6755	CG	LEU	A	131	9.176	94.291	14.219	1.00	31.94	T6
	ATOM	6756	CD1	LEU	A	131	9.947	93.121	14.760	1.00	24.97	T6
	ATOM	6757	CD2	LEU	A	131	7.775	94.282	14.770	1.00	31.09	T6
	ATOM	6758	C	LEU	A	131	11.610	97.268	13.998	1.00	29.96	T6
45	ATOM	6759	O	LEU	A	131	10.711	98.096	13.864	1.00	28.79	T6
	ATOM	6760	N	ASP	A	132	12.888	97.611	14.031	1.00	34.02	T6
	ATOM	6761	CA	ASP	A	132	13.280	99.003	13.903	1.00	30.59	T6
	ATOM	6762	CB	ASP	A	132	14.610	99.240	14.624	1.00	23.42	T6
	ATOM	6763	CG	ASP	A	132	14.436	99.469	16.111	1.00	29.39	T6
50	ATOM	6764	OD1	ASP	A	132	15.463	99.445	16.828	1.00	28.30	T6
	ATOM	6765	OD2	ASP	A	132	13.280	99.684	16.549	1.00	30.97	T6
	ATOM	6766	C	ASP	A	132	13.395	99.434	12.441	1.00	32.90	T6
	ATOM	6767	O	ASP	A	132	13.847	98.675	11.583	1.00	32.13	T6
	ATOM	6768	N	GLY	A	133	12.981	100.666	12.176	1.00	32.31	T6
55	ATOM	6769	CA	GLY	A	133	13.031	101.201	10.835	1.00	26.53	T6
	ATOM	6770	C	GLY	A	133	14.381	101.139	10.157	1.00	37.23	T6
	ATOM	6771	O	GLY	A	133	14.428	101.002	8.940	1.00	29.89	T6
	ATOM	6772	N	ASP	A	134	15.475	101.238	10.911	1.00	30.27	T6
	ATOM	6773	CA	ASP	A	134	16.802	101.189	10.300	1.00	35.37	T6
60	ATOM	6774	CB	ASP	A	134	17.923	101.506	11.280	1.00	32.74	T6
	ATOM	6775	CG	ASP	A	134	17.489	102.336	12.415	1.00	30.03	T6
	ATOM	6776	OD1	ASP	A	134	17.090	103.475	12.139	1.00	35.03	T6
	ATOM	6777	OD2	ASP	A	134	17.562	101.858	13.572	1.00	31.85	T6
	ATOM	6778	C	ASP	A	134	17.133	99.809	9.817	1.00	27.05	T6
65	ATOM	6779	O	ASP	A	134	17.432	99.594	8.660	1.00	29.93	T6
	ATOM	6780	N	VAL	A	135	17.108	98.876	10.754	1.00	27.19	T6

	ATOM	6781	CA	VAL	A	135	17.483	97.508	10.495	1.00	34.05	T6
	ATOM	6782	CB	VAL	A	135	17.693	96.804	11.820	1.00	29.32	T6
	ATOM	6783	CG1	VAL	A	135	18.793	97.515	12.587	1.00	31.29	T6
	ATOM	6784	CG2	VAL	A	135	16.412	96.821	12.619	1.00	31.99	T6
5	ATOM	6785	C	VAL	A	135	16.659	96.625	9.571	1.00	29.25	T6
	ATOM	6786	O	VAL	A	135	17.241	95.834	8.841	1.00	26.57	T6
	ATOM	6787	N	THR	A	136	15.334	96.721	9.583	1.00	27.78	T6
	ATOM	6788	CA	THR	A	136	14.570	95.862	8.677	1.00	29.89	T6
	ATOM	6789	CB	THR	A	136	13.824	94.737	9.453	1.00	24.67	T6
10	ATOM	6790	OG1	THR	A	136	12.661	95.268	10.078	1.00	32.29	T6
	ATOM	6791	CG2	THR	A	136	14.714	94.154	10.531	1.00	29.55	T6
	ATOM	6792	C	THR	A	136	13.596	96.629	7.766	1.00	33.78	T6
	ATOM	6793	O	THR	A	136	12.701	97.332	8.230	1.00	26.88	T6
	ATOM	6794	N	PHE	A	137	13.794	96.486	6.459	1.00	34.63	T6
15	ATOM	6795	CA	PHE	A	137	12.978	97.175	5.471	1.00	28.65	T6
	ATOM	6796	CB	PHE	A	137	13.625	98.521	5.143	1.00	27.57	T6
	ATOM	6797	CG	PHE	A	137	15.119	98.451	4.946	1.00	34.42	T6
	ATOM	6798	CD1	PHE	A	137	15.664	98.024	3.743	1.00	26.03	T6
	ATOM	6799	CD2	PHE	A	137	15.978	98.815	5.966	1.00	29.58	T6
20	ATOM	6800	CE1	PHE	A	137	17.032	97.968	3.569	1.00	34.34	T6
	ATOM	6801	CE2	PHE	A	137	17.347	98.757	5.793	1.00	37.28	T6
	ATOM	6802	CZ	PHE	A	137	17.872	98.335	4.598	1.00	34.44	T6
	ATOM	6803	C	PHE	A	137	12.774	96.346	4.198	1.00	32.53	T6
	ATOM	6804	O	PHE	A	137	13.495	95.379	3.967	1.00	24.33	T6
25	ATOM	6805	N	PHE	A	138	11.808	96.739	3.366	1.00	29.25	T6
	ATOM	6806	CA	PHE	A	138	11.488	95.995	2.152	1.00	34.98	T6
	ATOM	6807	CB	PHE	A	138	10.149	95.294	2.373	1.00	27.02	T6
	ATOM	6808	CG	PHE	A	138	9.834	94.231	1.374	1.00	33.23	T6
	ATOM	6809	CD1	PHE	A	138	10.838	93.607	0.653	1.00	33.38	T6
30	ATOM	6810	CD2	PHE	A	138	8.516	93.860	1.147	1.00	32.43	T6
	ATOM	6811	CE1	PHE	A	138	10.532	92.630	-0.291	1.00	32.10	T6
	ATOM	6812	CE2	PHE	A	138	8.198	92.885	0.207	1.00	32.18	T6
	ATOM	6813	CZ	PHE	A	138	9.207	92.270	-0.514	1.00	29.90	T6
	ATOM	6814	C	PHE	A	138	11.478	96.883	0.906	1.00	28.76	T6
35	ATOM	6815	O	PHE	A	138	10.854	97.938	0.878	1.00	31.66	T6
	ATOM	6816	N	GLY	A	139	12.170	96.399	-0.126	1.00	33.97	T6
	ATOM	6817	CA	GLY	A	139	12.388	97.114	-1.382	1.00	25.29	T6
	ATOM	6818	C	GLY	A	139	11.403	97.386	-2.489	1.00	29.54	T6
	ATOM	6819	O	GLY	A	139	10.313	97.840	-2.214	1.00	30.40	T6
40	ATOM	6820	N	ALA	A	140	11.829	97.168	-3.737	1.00	35.36	T6
	ATOM	6821	CA	ALA	A	140	11.029	97.383	-4.962	1.00	33.83	T6
	ATOM	6822	CB	ALA	A	140	9.574	97.028	-4.721	1.00	34.18	T6
	ATOM	6823	C	ALA	A	140	11.099	98.767	-5.626	1.00	24.35	T6
	ATOM	6824	O	ALA	A	140	10.499	99.732	-5.152	1.00	24.96	T6
45	ATOM	6825	N	LEU	A	141	11.818	98.835	-6.745	1.00	36.28	T6
	ATOM	6826	CA	LEU	A	141	11.993	100.062	-7.534	1.00	30.84	T6
	ATOM	6827	CB	LEU	A	141	13.350	100.705	-7.228	1.00	31.20	T6
	ATOM	6828	CG	LEU	A	141	13.835	101.891	-8.070	1.00	33.44	T6
	ATOM	6829	CD1	LEU	A	141	14.943	102.606	-7.357	1.00	26.45	T6
50	ATOM	6830	CD2	LEU	A	141	14.330	101.408	-9.397	1.00	29.33	T6
	ATOM	6831	C	LEU	A	141	11.932	99.679	-9.007	1.00	31.26	T6
	ATOM	6832	O	LEU	A	141	12.563	98.707	-9.423	1.00	27.50	T6
	ATOM	6833	N	LYS	A	142	11.194	100.444	-9.803	1.00	30.28	T6
	ATOM	6834	CA	LYS	A	142	11.058	100.122	-11.223	1.00	27.82	T6
55	ATOM	6835	CB	LYS	A	142	9.689	100.560	-11.737	1.00	24.74	T6
	ATOM	6836	CG	LYS	A	142	9.495	100.241	-13.197	1.00	33.62	T6
	ATOM	6837	CD	LYS	A	142	8.062	100.403	-13.645	1.00	31.82	T6
	ATOM	6838	CE	LYS	A	142	7.941	100.040	-15.118	1.00	34.37	T6
	ATOM	6839	NZ	LYS	A	142	6.552	100.200	-15.611	1.00	30.99	T6
60	ATOM	6840	C	LYS	A	142	12.136	100.692	-12.140	1.00	28.32	T6
	ATOM	6841	O	LYS	A	142	12.413	101.894	-12.116	1.00	25.55	T6
	ATOM	6842	N	LEU	A	143	12.734	99.822	-12.957	1.00	29.93	T6
	ATOM	6843	CA	LEU	A	143	13.783	100.232	-13.893	1.00	33.38	T6
	ATOM	6844	CB	LEU	A	143	14.650	99.035	-14.285	1.00	27.14	T6
65	ATOM	6845	CG	LEU	A	143	15.358	98.255	-13.179	1.00	29.54	T6
	ATOM	6846	CD1	LEU	A	143	16.067	97.063	-13.788	1.00	27.51	T6

	ATOM	6847	CD2	LEU	A	143	16.347	99.149	-12.467	1.00	25.30	T6
	ATOM	6848	C	LEU	A	143	13.163	100.822	-15.152	1.00	31.11	T6
	ATOM	6849	O	LEU	A	143	12.023	100.510	-15.490	1.00	39.97	T6
	ATOM	6850	N	LEU	A	144	13.912	101.675	-15.844	1.00	33.14	T6
5	ATOM	6851	CA	LEU	A	144	13.420	102.282	-17.075	1.00	24.36	T6
	ATOM	6852	CB	LEU	A	144	14.073	103.641	-17.304	1.00	24.76	T6
	ATOM	6853	CG	LEU	A	144	13.723	104.716	-16.281	1.00	30.66	T6
	ATOM	6854	CD1	LEU	A	144	14.580	105.937	-16.544	1.00	27.44	T6
	ATOM	6855	CD2	LEU	A	144	12.240	105.074	-16.362	1.00	31.00	T6
10	ATOM	6856	C	LEU	A	144	13.719	101.382	-18.264	1.00	30.64	T6
	ATOM	6857	O	LEU	A	144	14.577	100.485	-18.124	1.00	31.28	T6
	ATOM	6858	OXT	LEU	A	144	13.097	101.595	-19.325	1.00	37.85	T6
	ATOM	6859	CB	VAL	A	1	-33.125	107.279	-44.147	1.00	31.06	T7
	ATOM	6860	CG1	VAL	A	1	-33.999	108.464	-44.627	1.00	32.69	T7
15	ATOM	6861	CG2	VAL	A	1	-31.636	107.612	-44.335	1.00	30.25	T7
	ATOM	6862	C	VAL	A	1	-34.893	106.512	-42.500	1.00	32.56	T7
	ATOM	6863	O	VAL	A	1	-35.675	107.172	-41.802	1.00	24.93	T7
	ATOM	6864	N	VAL	A	1	-32.480	105.918	-42.105	1.00	32.59	T7
	ATOM	6865	CA	VAL	A	1	-33.420	106.953	-42.635	1.00	32.48	T7
20	ATOM	6866	N	THR	A	2	-35.277	105.413	-43.156	1.00	33.79	T7
	ATOM	6867	CA	THR	A	2	-36.663	104.943	-43.077	1.00	27.53	T7
	ATOM	6868	CB	THR	A	2	-37.282	104.762	-44.475	1.00	26.42	T7
	ATOM	6869	OG1	THR	A	2	-36.688	103.627	-45.116	1.00	28.09	T7
	ATOM	6870	CG2	THR	A	2	-37.052	106.000	-45.326	1.00	31.03	T7
25	ATOM	6871	C	THR	A	2	-36.801	103.615	-42.337	1.00	20.88	T7
	ATOM	6872	O	THR	A	2	-35.802	102.968	-42.008	1.00	22.57	T7
	ATOM	6873	N	GLN	A	3	-38.045	103.206	-42.094	1.00	30.87	T7
	ATOM	6874	CA	GLN	A	3	-38.321	101.949	-41.398	1.00	29.97	T7
	ATOM	6875	CB	GLN	A	3	-39.239	102.199	-40.209	1.00	39.04	T7
30	ATOM	6876	CG	GLN	A	3	-38.837	103.372	-39.345	1.00	35.68	T7
	ATOM	6877	CD	GLN	A	3	-39.692	103.474	-38.100	1.00	26.90	T7
	ATOM	6878	OE1	GLN	A	3	-39.707	102.557	-37.283	1.00	27.63	T7
	ATOM	6879	NE2	GLN	A	3	-40.410	104.586	-37.951	1.00	29.72	T7
	ATOM	6880	C	GLN	A	3	-38.988	100.928	-42.309	1.00	36.35	T7
35	ATOM	6881	O	GLN	A	3	-40.181	101.041	-42.602	1.00	26.99	T7
	ATOM	6882	N	ASP	A	4	-38.232	99.931	-42.753	1.00	34.12	T7
	ATOM	6883	CA	ASP	A	4	-38.810	98.910	-43.618	1.00	31.40	T7
	ATOM	6884	CB	ASP	A	4	-37.746	97.905	-44.066	1.00	34.13	T7
	ATOM	6885	CG	ASP	A	4	-36.654	98.543	-44.901	1.00	32.62	T7
40	ATOM	6886	OD1	ASP	A	4	-36.881	99.673	-45.402	1.00	34.22	T7
	ATOM	6887	OD2	ASP	A	4	-35.579	97.912	-45.065	1.00	33.41	T7
	ATOM	6888	C	ASP	A	4	-39.917	98.172	-42.883	1.00	26.23	T7
	ATOM	6889	O	ASP	A	4	-39.857	97.996	-41.670	1.00	34.01	T7
	ATOM	6890	N	CYS	A	5	-40.933	97.758	-43.629	1.00	32.20	T7
45	ATOM	6891	CA	CYS	A	5	-42.060	97.020	-43.078	1.00	31.40	T7
	ATOM	6892	CB	CYS	A	5	-43.085	97.977	-42.443	1.00	28.12	T7
	ATOM	6893	SG	CYS	A	5	-43.195	99.622	-43.193	1.00	27.08	T7
	ATOM	6894	C	CYS	A	5	-42.725	96.187	-44.168	1.00	36.63	T7
	ATOM	6895	O	CYS	A	5	-42.730	96.565	-45.327	1.00	26.21	T7
50	ATOM	6896	N	LEU	A	6	-43.253	95.032	-43.798	1.00	30.84	T7
	ATOM	6897	CA	LEU	A	6	-43.945	94.172	-44.745	1.00	30.30	T7
	ATOM	6898	CB	LEU	A	6	-43.047	93.023	-45.177	1.00	25.21	T7
	ATOM	6899	CG	LEU	A	6	-43.654	91.998	-46.136	1.00	32.64	T7
	ATOM	6900	CD1	LEU	A	6	-42.572	91.406	-47.004	1.00	37.63	T7
55	ATOM	6901	CD2	LEU	A	6	-44.348	90.914	-45.344	1.00	31.40	T7
	ATOM	6902	C	LEU	A	6	-45.187	93.640	-44.049	1.00	33.68	T7
	ATOM	6903	O	LEU	A	6	-45.114	93.204	-42.905	1.00	31.39	T7
	ATOM	6904	N	GLN	A	7	-46.327	93.677	-44.731	1.00	32.47	T7
	ATOM	6905	CA	GLN	A	7	-47.566	93.213	-44.130	1.00	33.21	T7
60	ATOM	6906	CB	GLN	A	7	-48.465	94.404	-43.833	1.00	29.68	T7
	ATOM	6907	CG	GLN	A	7	-49.657	94.085	-42.966	1.00	24.80	T7
	ATOM	6908	CD	GLN	A	7	-50.362	95.336	-42.506	1.00	25.27	T7
	ATOM	6909	OE1	GLN	A	7	-50.984	96.035	-43.292	1.00	30.31	T7
	ATOM	6910	NE2	GLN	A	7	-50.249	95.636	-41.230	1.00	28.49	T7
65	ATOM	6911	C	GLN	A	7	-48.302	92.229	-45.011	1.00	31.48	T7
	ATOM	6912	O	GLN	A	7	-48.342	92.385	-46.223	1.00	28.87	T7

	ATOM	6913	N	LEU	A	8	-48.884	91.213	-44.388	1.00	31.62	T7
	ATOM	6914	CA	LEU	A	8	-49.619	90.192	-45.105	1.00	35.80	T7
	ATOM	6915	CB	LEU	A	8	-49.012	88.814	-44.825	1.00	37.82	T7
	ATOM	6916	CG	LEU	A	8	-47.839	88.286	-45.663	1.00	27.54	T7
5	ATOM	6917	CD1	LEU	A	8	-47.202	89.387	-46.463	1.00	29.37	T7
	ATOM	6918	CD2	LEU	A	8	-46.839	87.637	-44.745	1.00	34.34	T7
	ATOM	6919	C	LEU	A	8	-51.091	90.197	-44.725	1.00	33.20	T7
	ATOM	6920	O	LEU	A	8	-51.462	90.607	-43.628	1.00	34.99	T7
	ATOM	6921	N	ILE	A	9	-51.920	89.727	-45.650	1.00	28.32	T7
10	ATOM	6922	CA	ILE	A	9	-53.370	89.654	-45.479	1.00	34.21	T7
	ATOM	6923	CB	ILE	A	9	-54.066	90.634	-46.432	1.00	27.83	T7
	ATOM	6924	CG2	ILE	A	9	-55.507	90.260	-46.609	1.00	35.14	T7
	ATOM	6925	CG1	ILE	A	9	-53.975	92.050	-45.903	1.00	26.37	T7
	ATOM	6926	CD1	ILE	A	9	-54.637	93.018	-46.843	1.00	27.69	T7
15	ATOM	6927	C	ILE	A	9	-53.866	88.251	-45.826	1.00	34.83	T7
	ATOM	6928	O	ILE	A	9	-53.351	87.620	-46.743	1.00	32.57	T7
	ATOM	6929	N	ALA	A	10	-54.877	87.767	-45.119	1.00	27.78	T7
	ATOM	6930	CA	ALA	A	10	-55.403	86.443	-45.420	1.00	31.36	T7
	ATOM	6931	CB	ALA	A	10	-56.484	86.064	-44.441	1.00	18.29	T7
20	ATOM	6932	C	ALA	A	10	-55.957	86.387	-46.835	1.00	25.90	T7
	ATOM	6933	O	ALA	A	10	-56.722	87.258	-47.255	1.00	28.17	T7
	ATOM	6934	N	ASP	A	11	-55.565	85.354	-47.573	1.00	31.80	T7
	ATOM	6935	CA	ASP	A	11	-56.041	85.171	-48.939	1.00	31.18	T7
	ATOM	6936	CB	ASP	A	11	-54.968	84.523	-49.808	1.00	35.98	T7
25	ATOM	6937	CG	ASP	A	11	-55.473	84.208	-51.195	1.00	25.84	T7
	ATOM	6938	OD1	ASP	A	11	-56.243	85.035	-51.730	1.00	31.76	T7
	ATOM	6939	OD2	ASP	A	11	-55.101	83.149	-51.750	1.00	29.22	T7
	ATOM	6940	C	ASP	A	11	-57.286	84.303	-48.929	1.00	32.66	T7
	ATOM	6941	O	ASP	A	11	-57.216	83.076	-49.002	1.00	30.38	T7
30	ATOM	6942	N	SER	A	12	-58.427	84.975	-48.834	1.00	31.86	T7
	ATOM	6943	CA	SER	A	12	-59.733	84.338	-48.784	1.00	33.85	T7
	ATOM	6944	CB	SER	A	12	-60.817	85.395	-48.536	1.00	30.79	T7
	ATOM	6945	OG	SER	A	12	-60.815	86.414	-49.534	1.00	30.85	T7
	ATOM	6946	C	SER	A	12	-60.079	83.565	-50.039	1.00	24.12	T7
35	ATOM	6947	O	SER	A	12	-61.254	83.323	-50.304	1.00	34.70	T7
	ATOM	6948	N	GLU	A	13	-59.073	83.179	-50.819	1.00	35.70	T7
	ATOM	6949	CA	GLU	A	13	-59.351	82.442	-52.045	1.00	28.17	T7
	ATOM	6950	CB	GLU	A	13	-59.322	83.381	-53.240	1.00	29.93	T7
	ATOM	6951	CG	GLU	A	13	-60.671	83.978	-53.503	1.00	31.47	T7
40	ATOM	6952	CD	GLU	A	13	-60.664	84.831	-54.742	1.00	39.45	T7
	ATOM	6953	OE1	GLU	A	13	-60.032	84.396	-55.745	1.00	25.41	T7
	ATOM	6954	OE2	GLU	A	13	-61.288	85.930	-54.725	1.00	25.39	T7
	ATOM	6955	C	GLU	A	13	-58.492	81.221	-52.314	1.00	25.96	T7
	ATOM	6956	O	GLU	A	13	-58.323	80.802	-53.459	1.00	29.71	T7
45	ATOM	6957	N	THR	A	14	-57.953	80.657	-51.246	1.00	31.48	T7
	ATOM	6958	CA	THR	A	14	-57.165	79.449	-51.347	1.00	35.17	T7
	ATOM	6959	CB	THR	A	14	-55.659	79.726	-51.540	1.00	23.06	T7
	ATOM	6960	OG1	THR	A	14	-55.181	80.521	-50.458	1.00	27.18	T7
	ATOM	6961	CG2	THR	A	14	-55.399	80.448	-52.859	1.00	28.75	T7
50	ATOM	6962	C	THR	A	14	-57.394	78.726	-50.037	1.00	35.12	T7
	ATOM	6963	O	THR	A	14	-57.658	79.344	-49.011	1.00	30.59	T7
	ATOM	6964	N	PRO	A	15	-57.319	77.398	-50.061	1.00	30.05	T7
	ATOM	6965	CD	PRO	A	15	-56.996	76.569	-51.232	1.00	24.97	T7
	ATOM	6966	CA	PRO	A	15	-57.526	76.572	-48.872	1.00	31.85	T7
55	ATOM	6967	CB	PRO	A	15	-57.383	75.148	-49.409	1.00	30.66	T7
	ATOM	6968	CG	PRO	A	15	-57.682	75.284	-50.889	1.00	23.47	T7
	ATOM	6969	C	PRO	A	15	-56.493	76.869	-47.808	1.00	32.10	T7
	ATOM	6970	O	PRO	A	15	-55.353	77.173	-48.118	1.00	37.01	T7
	ATOM	6971	N	THR	A	16	-56.895	76.777	-46.553	1.00	29.21	T7
60	ATOM	6972	CA	THR	A	16	-55.964	77.014	-45.463	1.00	31.32	T7
	ATOM	6973	CB	THR	A	16	-56.711	77.110	-44.146	1.00	24.10	T7
	ATOM	6974	OG1	THR	A	16	-57.215	75.818	-43.778	1.00	27.54	T7
	ATOM	6975	CG2	THR	A	16	-57.875	78.071	-44.303	1.00	28.45	T7
	ATOM	6976	C	THR	A	16	-54.989	75.843	-45.389	1.00	27.55	T7
65	ATOM	6977	O	THR	A	16	-55.397	74.715	-45.108	1.00	32.14	T7
	ATOM	6978	N	ILE	A	17	-53.708	76.107	-45.626	1.00	33.38	T7

	ATOM	6979	CA	ILE	A	17	-52.693	75.055	-45.611	1.00	35.78	T7
	ATOM	6980	CB	ILE	A	17	-51.302	75.656	-45.769	1.00	31.70	T7
	ATOM	6981	CG2	ILE	A	17	-50.258	74.562	-45.726	1.00	30.92	T7
	ATOM	6982	CG1	ILE	A	17	-51.226	76.429	-47.085	1.00	31.77	T7
5	ATOM	6983	CD1	ILE	A	17	-49.912	77.127	-47.307	1.00	32.89	T7
	ATOM	6984	C	ILE	A	17	-52.691	74.160	-44.372	1.00	25.24	T7
	ATOM	6985	O	ILE	A	17	-52.729	74.645	-43.244	1.00	29.92	T7
	ATOM	6986	N	GLN	A	18	-52.651	72.848	-44.597	1.00	35.37	T7
	ATOM	6987	CA	GLN	A	18	-52.631	71.871	-43.509	1.00	27.00	T7
10	ATOM	6988	CB	GLN	A	18	-53.770	70.874	-43.663	1.00	29.69	T7
	ATOM	6989	CG	GLN	A	18	-54.639	70.808	-42.438	1.00	27.52	T7
	ATOM	6990	CD	GLN	A	18	-55.511	72.030	-42.321	1.00	27.41	T7
	ATOM	6991	OE1	GLN	A	18	-56.509	72.160	-43.029	1.00	35.50	T7
	ATOM	6992	NE2	GLN	A	18	-55.132	72.949	-41.439	1.00	34.68	T7
15	ATOM	6993	C	GLN	A	18	-51.307	71.111	-43.483	1.00	28.18	T7
	ATOM	6994	O	GLN	A	18	-50.821	70.656	-44.518	1.00	32.10	T7
	ATOM	6995	N	LYS	A	19	-50.728	70.957	-42.298	1.00	24.54	T7
	ATOM	6996	CA	LYS	A	19	-49.447	70.271	-42.188	1.00	24.60	T7
	ATOM	6997	CB	LYS	A	19	-48.365	71.085	-42.900	1.00	32.26	T7
20	ATOM	6998	CG	LYS	A	19	-46.961	70.598	-42.631	1.00	25.92	T7
	ATOM	6999	CD	LYS	A	19	-45.956	71.291	-43.534	1.00	37.67	T7
	ATOM	7000	CE	LYS	A	19	-44.524	70.806	-43.260	1.00	28.78	T7
	ATOM	7001	NZ	LYS	A	19	-43.512	71.440	-44.167	1.00	32.70	T7
	ATOM	7002	C	LYS	A	19	-49.030	70.024	-40.743	1.00	33.36	T7
25	ATOM	7003	O	LYS	A	19	-49.030	70.949	-39.927	1.00	26.67	T7
	ATOM	7004	N	GLY	A	20	-48.660	68.780	-40.439	1.00	28.11	T7
	ATOM	7005	CA	GLY	A	20	-48.245	68.429	-39.092	1.00	32.86	T7
	ATOM	7006	C	GLY	A	20	-49.321	68.796	-38.095	1.00	30.40	T7
	ATOM	7007	O	GLY	A	20	-49.023	69.297	-37.011	1.00	31.95	T7
30	ATOM	7008	N	SER	A	21	-50.574	68.547	-38.475	1.00	28.36	T7
	ATOM	7009	CA	SER	A	21	-51.740	68.861	-37.641	1.00	35.64	T7
	ATOM	7010	CB	SER	A	21	-51.843	67.880	-36.463	1.00	36.69	T7
	ATOM	7011	OG	SER	A	21	-50.706	67.951	-35.629	1.00	27.22	T7
	ATOM	7012	C	SER	A	21	-51.734	70.317	-37.134	1.00	32.28	T7
35	ATOM	7013	O	SER	A	21	-52.098	70.605	-35.986	1.00	29.37	T7
	ATOM	7014	N	TYR	A	22	-51.316	71.219	-38.023	1.00	30.04	T7
	ATOM	7015	CA	TYR	A	22	-51.251	72.651	-37.769	1.00	32.62	T7
	ATOM	7016	CB	TYR	A	22	-49.802	73.122	-37.750	1.00	30.68	T7
	ATOM	7017	CG	TYR	A	22	-49.191	73.201	-36.379	1.00	29.60	T7
40	ATOM	7018	CD1	TYR	A	22	-49.669	72.411	-35.333	1.00	30.81	T7
	ATOM	7019	CE1	TYR	A	22	-49.082	72.455	-34.063	1.00	38.04	T7
	ATOM	7020	CD2	TYR	A	22	-48.107	74.040	-36.126	1.00	34.27	T7
	ATOM	7021	CE2	TYR	A	22	-47.509	74.088	-34.861	1.00	27.64	T7
	ATOM	7022	CZ	TYR	A	22	-48.005	73.292	-33.836	1.00	34.76	T7
45	ATOM	7023	OH	TYR	A	22	-47.432	73.329	-32.586	1.00	34.99	T7
	ATOM	7024	C	TYR	A	22	-51.963	73.317	-38.928	1.00	28.14	T7
	ATOM	7025	O	TYR	A	22	-51.919	72.810	-40.048	1.00	29.24	T7
	ATOM	7026	N	THR	A	23	-52.630	74.435	-38.671	1.00	30.62	T7
	ATOM	7027	CA	THR	A	23	-53.308	75.137	-39.746	1.00	31.91	T7
50	ATOM	7028	CB	THR	A	23	-54.751	75.512	-39.380	1.00	33.73	T7
	ATOM	7029	OG1	THR	A	23	-55.379	74.433	-38.678	1.00	23.57	T7
	ATOM	7030	CG2	THR	A	23	-55.541	75.777	-40.634	1.00	27.53	T7
	ATOM	7031	C	THR	A	23	-52.528	76.405	-40.044	1.00	35.27	T7
	ATOM	7032	O	THR	A	23	-52.174	77.153	-39.134	1.00	32.98	T7
55	ATOM	7033	N	PHE	A	24	-52.240	76.631	-41.320	1.00	25.34	T7
	ATOM	7034	CA	PHE	A	24	-51.503	77.812	-41.744	1.00	27.76	T7
	ATOM	7035	CB	PHE	A	24	-50.210	77.406	-42.438	1.00	30.71	T7
	ATOM	7036	CG	PHE	A	24	-49.233	76.724	-41.537	1.00	25.16	T7
	ATOM	7037	CD1	PHE	A	24	-49.370	75.378	-41.231	1.00	33.77	T7
60	ATOM	7038	CD2	PHE	A	24	-48.184	77.441	-40.970	1.00	27.58	T7
	ATOM	7039	CE1	PHE	A	24	-48.475	74.751	-40.371	1.00	38.39	T7
	ATOM	7040	CE2	PHE	A	24	-47.287	76.825	-40.111	1.00	23.22	T7
	ATOM	7041	CZ	PHE	A	24	-47.434	75.474	-39.811	1.00	28.84	T7
	ATOM	7042	C	PHE	A	24	-52.323	78.700	-42.671	1.00	32.62	T7
65	ATOM	7043	O	PHE	A	24	-52.803	78.255	-43.708	1.00	30.13	T7
	ATOM	7044	N	VAL	A	25	-52.473	79.959	-42.282	1.00	29.37	T7

	ATOM	7045	CA	VAL	A	25	-53.230	80.927	-43.052	1.00	35.46	T7
	ATOM	7046	CB	VAL	A	25	-53.348	82.267	-42.301	1.00	34.93	T7
	ATOM	7047	CG1	VAL	A	25	-54.058	83.284	-43.155	1.00	28.98	T7
	ATOM	7048	CG2	VAL	A	25	-54.092	82.069	-41.009	1.00	33.57	T7
5	ATOM	7049	C	VAL	A	25	-52.554	81.196	-44.383	1.00	29.39	T7
	ATOM	7050	O	VAL	A	25	-51.343	81.389	-44.441	1.00	29.66	T7
	ATOM	7051	N	PRO	A	26	-53.332	81.189	-45.477	1.00	29.37	T7
	ATOM	7052	CD	PRO	A	26	-54.739	80.757	-45.538	1.00	38.85	T7
	ATOM	7053	CA	PRO	A	26	-52.808	81.444	-46.825	1.00	27.28	T7
10	ATOM	7054	CB	PRO	A	26	-53.945	80.982	-47.734	1.00	35.89	T7
	ATOM	7055	CG	PRO	A	26	-54.783	80.078	-46.861	1.00	24.09	T7
	ATOM	7056	C	PRO	A	26	-52.595	82.955	-46.933	1.00	25.73	T7
	ATOM	7057	O	PRO	A	26	-53.562	83.716	-46.942	1.00	33.45	T7
	ATOM	7058	N	TRP	A	27	-51.351	83.402	-47.018	1.00	35.80	T7
15	ATOM	7059	CA	TRP	A	27	-51.105	84.835	-47.092	1.00	39.72	T7
	ATOM	7060	CB	TRP	A	27	-49.768	85.181	-46.431	1.00	29.60	T7
	ATOM	7061	CG	TRP	A	27	-49.740	84.877	-44.976	1.00	31.26	T7
	ATOM	7062	CD2	TRP	A	27	-50.672	85.328	-43.986	1.00	34.23	T7
	ATOM	7063	CE2	TRP	A	27	-50.295	84.739	-42.761	1.00	38.64	T7
20	ATOM	7064	CE3	TRP	A	27	-51.788	86.168	-44.016	1.00	35.24	T7
	ATOM	7065	CD1	TRP	A	27	-48.857	84.064	-44.330	1.00	32.80	T7
	ATOM	7066	NE1	TRP	A	27	-49.184	83.973	-42.999	1.00	27.26	T7
	ATOM	7067	CZ2	TRP	A	27	-50.998	84.963	-41.574	1.00	29.78	T7
	ATOM	7068	CZ3	TRP	A	27	-52.485	86.389	-42.835	1.00	25.46	T7
25	ATOM	7069	CH2	TRP	A	27	-52.085	85.786	-41.633	1.00	32.59	T7
	ATOM	7070	C	TRP	A	27	-51.149	85.460	-48.478	1.00	37.67	T7
	ATOM	7071	O	TRP	A	27	-51.087	84.783	-49.504	1.00	27.76	T7
	ATOM	7072	N	LEU	A	28	-51.263	86.780	-48.478	1.00	31.56	T7
	ATOM	7073	CA	LEU	A	28	-51.306	87.568	-49.692	1.00	30.13	T7
30	ATOM	7074	CB	LEU	A	28	-52.748	87.752	-50.133	1.00	39.14	T7
	ATOM	7075	CG	LEU	A	28	-52.928	88.137	-51.595	1.00	35.82	T7
	ATOM	7076	CD1	LEU	A	28	-52.393	87.001	-52.479	1.00	30.15	T7
	ATOM	7077	CD2	LEU	A	28	-54.411	88.409	-51.865	1.00	22.50	T7
	ATOM	7078	C	LEU	A	28	-50.688	88.914	-49.334	1.00	32.63	T7
35	ATOM	7079	O	LEU	A	28	-51.072	89.529	-48.347	1.00	30.16	T7
	ATOM	7080	N	LEU	A	29	-49.727	89.369	-50.123	1.00	29.67	T7
	ATOM	7081	CA	LEU	A	29	-49.069	90.623	-49.821	1.00	30.83	T7
	ATOM	7082	CB	LEU	A	29	-48.056	90.969	-50.902	1.00	27.33	T7
	ATOM	7083	CG	LEU	A	29	-47.356	92.298	-50.636	1.00	29.28	T7
40	ATOM	7084	CD1	LEU	A	29	-46.332	92.126	-49.536	1.00	26.39	T7
	ATOM	7085	CD2	LEU	A	29	-46.695	92.781	-51.892	1.00	25.59	T7
	ATOM	7086	C	LEU	A	29	-50.028	91.782	-49.676	1.00	24.57	T7
	ATOM	7087	O	LEU	A	29	-50.831	92.040	-50.561	1.00	25.59	T7
	ATOM	7088	N	SER	A	30	-49.946	92.473	-48.545	1.00	29.66	T7
45	ATOM	7089	CA	SER	A	30	-50.774	93.641	-48.316	1.00	30.59	T7
	ATOM	7090	CB	SER	A	30	-51.007	93.864	-46.832	1.00	31.79	T7
	ATOM	7091	OG	SER	A	30	-51.709	95.073	-46.630	1.00	24.45	T7
	ATOM	7092	C	SER	A	30	-49.956	94.787	-48.886	1.00	29.67	T7
	ATOM	7093	O	SER	A	30	-50.421	95.539	-49.735	1.00	34.98	T7
50	ATOM	7094	N	PHE	A	31	-48.723	94.907	-48.413	1.00	31.63	T7
	ATOM	7095	CA	PHE	A	31	-47.819	95.939	-48.892	1.00	38.60	T7
	ATOM	7096	CB	PHE	A	31	-48.265	97.320	-48.401	1.00	25.52	T7
	ATOM	7097	CG	PHE	A	31	-47.744	97.686	-47.045	1.00	30.33	T7
	ATOM	7098	CD1	PHE	A	31	-46.486	98.258	-46.903	1.00	37.46	T7
55	ATOM	7099	CD2	PHE	A	31	-48.506	97.454	-45.905	1.00	25.24	T7
	ATOM	7100	CE1	PHE	A	31	-45.991	98.593	-45.650	1.00	27.49	T7
	ATOM	7101	CE2	PHE	A	31	-48.021	97.786	-44.647	1.00	36.62	T7
	ATOM	7102	CZ	PHE	A	31	-46.760	98.357	-44.518	1.00	26.10	T7
	ATOM	7103	C	PHE	A	31	-46.419	95.624	-48.395	1.00	23.72	T7
60	ATOM	7104	O	PHE	A	31	-46.247	94.962	-47.373	1.00	30.32	T7
	ATOM	7105	N	LYS	A	32	-45.419	96.087	-49.133	1.00	29.52	T7
	ATOM	7106	CA	LYS	A	32	-44.032	95.863	-48.765	1.00	28.94	T7
	ATOM	7107	CB	LYS	A	32	-43.413	94.796	-49.658	1.00	35.50	T7
	ATOM	7108	CG	LYS	A	32	-41.917	94.724	-49.533	1.00	32.52	T7
65	ATOM	7109	CD	LYS	A	32	-41.295	93.813	-50.554	1.00	39.10	T7
	ATOM	7110	CE	LYS	A	32	-39.797	94.006	-50.528	1.00	32.87	T7

	ATOM	7111	NZ	LYS	A	32	-39.117	93.027	-51.396	1.00	35.10	T7
	ATOM	7112	C	LYS	A	32	-43.274	97.170	-48.925	1.00	30.92	T7
	ATOM	7113	O	LYS	A	32	-43.314	97.793	-49.982	1.00	36.03	T7
	ATOM	7114	N	ARG	A	33	-42.581	97.589	-47.878	1.00	36.62	T7
5	ATOM	7115	CA	ARG	A	33	-41.843	98.840	-47.914	1.00	32.53	T7
	ATOM	7116	CB	ARG	A	33	-42.537	99.846	-46.997	1.00	28.52	T7
	ATOM	7117	CG	ARG	A	33	-41.830	101.163	-46.771	1.00	37.21	T7
	ATOM	7118	CD	ARG	A	33	-42.850	102.195	-46.289	1.00	36.40	T7
	ATOM	7119	NE	ARG	A	33	-42.254	103.480	-45.946	1.00	27.80	T7
10	ATOM	7120	CZ	ARG	A	33	-41.661	103.735	-44.785	1.00	33.02	T7
	ATOM	7121	NH1	ARG	A	33	-41.600	102.784	-43.854	1.00	34.61	T7
	ATOM	7122	NH2	ARG	A	33	-41.116	104.929	-44.561	1.00	31.30	T7
	ATOM	7123	C	ARG	A	33	-40.411	98.607	-47.475	1.00	22.30	T7
	ATOM	7124	O	ARG	A	33	-40.163	98.087	-46.385	1.00	28.88	T7
15	ATOM	7125	N	GLY	A	34	-39.464	98.974	-48.331	1.00	31.54	T7
	ATOM	7126	CA	GLY	A	34	-38.069	98.793	-47.985	1.00	34.68	T7
	ATOM	7127	C	GLY	A	34	-37.486	97.489	-48.475	1.00	31.27	T7
	ATOM	7128	O	GLY	A	34	-38.071	96.805	-49.313	1.00	34.22	T7
	ATOM	7129	N	SER	A	35	-36.337	97.128	-47.924	1.00	33.79	T7
20	ATOM	7130	CA	SER	A	35	-35.647	95.917	-48.335	1.00	27.44	T7
	ATOM	7131	CB	SER	A	35	-34.296	96.300	-48.923	1.00	29.59	T7
	ATOM	7132	OG	SER	A	35	-33.556	97.077	-47.983	1.00	37.14	T7
	ATOM	7133	C	SER	A	35	-35.415	94.872	-47.243	1.00	34.64	T7
	ATOM	7134	O	SER	A	35	-35.178	93.704	-47.545	1.00	22.96	T7
25	ATOM	7135	N	ALA	A	36	-35.481	95.279	-45.982	1.00	25.57	T7
	ATOM	7136	CA	ALA	A	36	-35.219	94.358	-44.887	1.00	24.07	T7
	ATOM	7137	CB	ALA	A	36	-35.189	95.118	-43.581	1.00	23.64	T7
	ATOM	7138	C	ALA	A	36	-36.144	93.161	-44.758	1.00	27.33	T7
	ATOM	7139	O	ALA	A	36	-35.799	92.202	-44.064	1.00	33.25	T7
30	ATOM	7140	N	LEU	A	37	-37.306	93.196	-45.410	1.00	30.09	T7
	ATOM	7141	CA	LEU	A	37	-38.254	92.082	-45.314	1.00	31.45	T7
	ATOM	7142	CB	LEU	A	37	-39.367	92.444	-44.338	1.00	30.57	T7
	ATOM	7143	CG	LEU	A	37	-38.897	92.750	-42.912	1.00	26.36	T7
	ATOM	7144	CD1	LEU	A	37	-39.950	93.532	-42.155	1.00	30.50	T7
35	ATOM	7145	CD2	LEU	A	37	-38.575	91.449	-42.207	1.00	28.19	T7
	ATOM	7146	C	LEU	A	37	-38.851	91.685	-46.652	1.00	35.38	T7
	ATOM	7147	O	LEU	A	37	-39.064	92.531	-47.514	1.00	35.53	T7
	ATOM	7148	N	GLU	A	38	-39.132	90.393	-46.806	1.00	31.90	T7
	ATOM	7149	CA	GLU	A	38	-39.685	89.844	-48.050	1.00	28.40	T7
40	ATOM	7150	CB	GLU	A	38	-38.562	89.310	-48.937	1.00	35.25	T7
	ATOM	7151	CG	GLU	A	38	-37.831	90.334	-49.771	1.00	25.28	T7
	ATOM	7152	CD	GLU	A	38	-36.571	89.757	-50.418	1.00	21.19	T7
	ATOM	7153	OE1	GLU	A	38	-36.594	88.560	-50.804	1.00	30.25	T7
	ATOM	7154	OE2	GLU	A	38	-35.565	90.505	-50.543	1.00	29.41	T7
45	ATOM	7155	C	GLU	A	38	-40.646	88.687	-47.811	1.00	32.18	T7
	ATOM	7156	O	GLU	A	38	-40.656	88.102	-46.731	1.00	29.38	T7
	ATOM	7157	N	GLU	A	39	-41.443	88.350	-48.825	1.00	23.02	T7
	ATOM	7158	CA	GLU	A	39	-42.364	87.221	-48.709	1.00	34.71	T7
	ATOM	7159	CB	GLU	A	39	-43.584	87.361	-49.599	1.00	28.18	T7
50	ATOM	7160	CG	GLU	A	39	-44.215	88.700	-49.640	1.00	29.18	T7
	ATOM	7161	CD	GLU	A	39	-44.880	88.933	-50.986	1.00	25.92	T7
	ATOM	7162	OE1	GLU	A	39	-44.244	89.584	-51.866	1.00	27.16	T7
	ATOM	7163	OE2	GLU	A	39	-46.026	88.440	-51.167	1.00	29.56	T7
	ATOM	7164	C	GLU	A	39	-41.600	86.041	-49.252	1.00	27.03	T7
55	ATOM	7165	O	GLU	A	39	-40.810	86.184	-50.179	1.00	31.79	T7
	ATOM	7166	N	LYS	A	40	-41.845	84.870	-48.696	1.00	25.32	T7
	ATOM	7167	CA	LYS	A	40	-41.173	83.688	-49.181	1.00	28.60	T7
	ATOM	7168	CB	LYS	A	40	-39.735	83.625	-48.673	1.00	28.08	T7
	ATOM	7169	CG	LYS	A	40	-38.980	82.416	-49.213	1.00	27.75	T7
60	ATOM	7170	CD	LYS	A	40	-37.714	82.146	-48.422	1.00	23.88	T7
	ATOM	7171	CE	LYS	A	40	-37.087	80.816	-48.825	1.00	30.07	T7
	ATOM	7172	NZ	LYS	A	40	-35.925	80.483	-47.950	1.00	33.28	T7
	ATOM	7173	C	LYS	A	40	-41.923	82.450	-48.739	1.00	25.81	T7
	ATOM	7174	O	LYS	A	40	-41.852	82.042	-47.578	1.00	36.64	T7
65	ATOM	7175	N	GLU	A	41	-42.659	81.866	-49.676	1.00	27.84	T7
	ATOM	7176	CA	GLU	A	41	-43.419	80.659	-49.408	1.00	32.42	T7

	ATOM	7177	CB	GLU	A	41	-42.447	79.489	-49.289	1.00	32.43	T7
	ATOM	7178	CG	GLU	A	41	-41.499	79.442	-50.471	1.00	27.68	T7
	ATOM	7179	CD	GLU	A	41	-40.300	78.546	-50.237	1.00	23.07	T7
	ATOM	7180	OE1	GLU	A	41	-39.614	78.708	-49.188	1.00	28.94	T7
5	ATOM	7181	OE2	GLU	A	41	-40.036	77.685	-51.114	1.00	32.19	T7
	ATOM	7182	C	GLU	A	41	-44.272	80.800	-48.150	1.00	26.41	T7
	ATOM	7183	O	GLU	A	41	-44.125	80.050	-47.186	1.00	30.69	T7
	ATOM	7184	N	ASN	A	42	-45.150	81.790	-48.171	1.00	34.48	T7
	ATOM	7185	CA	ASN	A	42	-46.062	82.035	-47.076	1.00	31.36	T7
10	ATOM	7186	CB	ASN	A	42	-46.961	80.829	-46.886	1.00	30.54	T7
	ATOM	7187	CG	ASN	A	42	-48.390	81.223	-46.644	1.00	31.48	T7
	ATOM	7188	OD1	ASN	A	42	-49.003	80.807	-45.660	1.00	27.30	T7
	ATOM	7189	ND2	ASN	A	42	-48.939	82.036	-47.544	1.00	31.09	T7
	ATOM	7190	C	ASN	A	42	-45.419	82.390	-45.751	1.00	31.54	T7
15	ATOM	7191	O	ASN	A	42	-46.042	82.257	-44.700	1.00	29.15	T7
	ATOM	7192	N	LYS	A	43	-44.177	82.846	-45.800	1.00	32.69	T7
	ATOM	7193	CA	LYS	A	43	-43.463	83.238	-44.596	1.00	36.53	T7
	ATOM	7194	CB	LYS	A	43	-42.418	82.187	-44.235	1.00	25.03	T7
	ATOM	7195	CG	LYS	A	43	-42.995	80.886	-43.747	1.00	26.52	T7
20	ATOM	7196	CD	LYS	A	43	-41.902	79.877	-43.474	1.00	27.16	T7
	ATOM	7197	CE	LYS	A	43	-41.274	79.387	-44.772	1.00	37.04	T7
	ATOM	7198	NZ	LYS	A	43	-40.229	78.340	-44.542	1.00	35.21	T7
	ATOM	7199	C	LYS	A	43	-42.779	84.569	-44.841	1.00	31.31	T7
	ATOM	7200	O	LYS	A	43	-42.601	84.972	-45.983	1.00	32.81	T7
25	ATOM	7201	N	ILE	A	44	-42.410	85.261	-43.773	1.00	30.71	T7
	ATOM	7202	CA	ILE	A	44	-41.726	86.531	-43.923	1.00	33.03	T7
	ATOM	7203	CB	ILE	A	44	-42.164	87.535	-42.861	1.00	32.12	T7
	ATOM	7204	CG2	ILE	A	44	-41.414	88.837	-43.049	1.00	31.90	T7
	ATOM	7205	CG1	ILE	A	44	-43.666	87.777	-42.980	1.00	28.61	T7
30	ATOM	7206	CD1	ILE	A	44	-44.217	88.735	-41.970	1.00	22.63	T7
	ATOM	7207	C	ILE	A	44	-40.239	86.273	-43.777	1.00	32.16	T7
	ATOM	7208	O	ILE	A	44	-39.797	85.720	-42.771	1.00	30.79	T7
	ATOM	7209	N	LEU	A	45	-39.468	86.666	-44.784	1.00	30.25	T7
	ATOM	7210	CA	LEU	A	45	-38.032	86.455	-44.765	1.00	32.51	T7
35	ATOM	7211	CB	LEU	A	45	-37.569	86.016	-46.144	1.00	34.45	T7
	ATOM	7212	CG	LEU	A	45	-36.060	85.841	-46.265	1.00	22.87	T7
	ATOM	7213	CD1	LEU	A	45	-35.621	84.613	-45.481	1.00	28.67	T7
	ATOM	7214	CD2	LEU	A	45	-35.695	85.711	-47.731	1.00	30.17	T7
	ATOM	7215	C	LEU	A	45	-37.248	87.691	-44.340	1.00	27.53	T7
40	ATOM	7216	O	LEU	A	45	-37.447	88.780	-44.883	1.00	29.73	T7
	ATOM	7217	N	VAL	A	46	-36.350	87.517	-43.372	1.00	30.75	T7
	ATOM	7218	CA	VAL	A	46	-35.538	88.623	-42.878	1.00	27.43	T7
	ATOM	7219	CB	VAL	A	46	-35.097	88.369	-41.437	1.00	38.31	T7
	ATOM	7220	CG1	VAL	A	46	-34.250	89.520	-40.947	1.00	31.83	T7
45	ATOM	7221	CG2	VAL	A	46	-36.300	88.200	-40.560	1.00	27.56	T7
	ATOM	7222	C	VAL	A	46	-34.295	88.820	-43.744	1.00	30.03	T7
	ATOM	7223	O	VAL	A	46	-33.461	87.921	-43.856	1.00	28.64	T7
	ATOM	7224	N	LYS	A	47	-34.157	89.997	-44.343	1.00	33.59	T7
	ATOM	7225	CA	LYS	A	47	-33.010	90.268	-45.205	1.00	29.40	T7
50	ATOM	7226	CB	LYS	A	47	-33.475	90.945	-46.500	1.00	30.65	T7
	ATOM	7227	CG	LYS	A	47	-34.136	90.010	-47.495	1.00	36.05	T7
	ATOM	7228	CD	LYS	A	47	-33.222	88.837	-47.788	1.00	28.92	T7
	ATOM	7229	CE	LYS	A	47	-33.674	88.041	-48.996	1.00	37.20	T7
	ATOM	7230	NZ	LYS	A	47	-33.434	88.760	-50.287	1.00	35.30	T7
55	ATOM	7231	C	LYS	A	47	-31.908	91.103	-44.566	1.00	36.50	T7
	ATOM	7232	O	LYS	A	47	-30.840	91.259	-45.135	1.00	29.84	T7
	ATOM	7233	N	GLU	A	48	-32.171	91.644	-43.389	1.00	32.41	T7
	ATOM	7234	CA	GLU	A	48	-31.201	92.465	-42.678	1.00	32.24	T7
	ATOM	7235	CB	GLU	A	48	-31.463	93.943	-42.902	1.00	24.18	T7
60	ATOM	7236	CG	GLU	A	48	-31.175	94.441	-44.285	1.00	28.39	T7
	ATOM	7237	CD	GLU	A	48	-31.591	95.895	-44.454	1.00	33.63	T7
	ATOM	7238	OE1	GLU	A	48	-31.367	96.696	-43.500	1.00	26.72	T7
	ATOM	7239	OE2	GLU	A	48	-32.134	96.229	-45.541	1.00	33.84	T7
	ATOM	7240	C	GLU	A	48	-31.387	92.199	-41.210	1.00	31.85	T7
65	ATOM	7241	O	GLU	A	48	-32.490	92.354	-40.693	1.00	30.13	T7
	ATOM	7242	N	THR	A	49	-30.328	91.816	-40.515	1.00	28.32	T7

	ATOM	7243	CA	THR	A	49	-30.486	91.550	-39.097	1.00	32.50	T7
	ATOM	7244	CB	THR	A	49	-29.245	90.869	-38.521	1.00	31.31	T7
	ATOM	7245	OG1	THR	A	49	-28.348	91.864	-38.035	1.00	34.41	T7
	ATOM	7246	CG2	THR	A	49	-28.549	90.046	-39.591	1.00	35.33	T7
5	ATOM	7247	C	THR	A	49	-30.756	92.868	-38.361	1.00	29.16	T7
	ATOM	7248	O	THR	A	49	-30.311	93.938	-38.792	1.00	32.18	T7
	ATOM	7249	N	GLY	A	50	-31.510	92.782	-37.267	1.00	19.78	T7
	ATOM	7250	CA	GLY	A	50	-31.832	93.962	-36.488	1.00	28.38	T7
	ATOM	7251	C	GLY	A	50	-32.976	93.672	-35.545	1.00	27.68	T7
10	ATOM	7252	O	GLY	A	50	-33.306	92.511	-35.309	1.00	35.44	T7
	ATOM	7253	N	TYR	A	51	-33.581	94.726	-35.003	1.00	28.02	T7
	ATOM	7254	CA	TYR	A	51	-34.714	94.585	-34.093	1.00	34.94	T7
	ATOM	7255	CB	TYR	A	51	-34.565	95.524	-32.897	1.00	28.19	T7
	ATOM	7256	CG	TYR	A	51	-33.522	95.045	-31.928	1.00	25.32	T7
15	ATOM	7257	CD1	TYR	A	51	-32.167	95.246	-32.173	1.00	29.99	T7
	ATOM	7258	CE1	TYR	A	51	-31.197	94.700	-31.335	1.00	27.63	T7
	ATOM	7259	CD2	TYR	A	51	-33.889	94.295	-30.815	1.00	24.26	T7
	ATOM	7260	CE2	TYR	A	51	-32.935	93.740	-29.970	1.00	35.30	T7
	ATOM	7261	CZ	TYR	A	51	-31.588	93.938	-30.232	1.00	29.84	T7
20	ATOM	7262	OH	TYR	A	51	-30.647	93.328	-29.413	1.00	36.27	T7
	ATOM	7263	C	TYR	A	51	-36.019	94.870	-34.818	1.00	27.67	T7
	ATOM	7264	O	TYR	A	51	-36.170	95.906	-35.468	1.00	36.23	T7
	ATOM	7265	N	PHE	A	52	-36.963	93.940	-34.704	1.00	30.79	T7
	ATOM	7266	CA	PHE	A	52	-38.248	94.087	-35.368	1.00	27.50	T7
25	ATOM	7267	CB	PHE	A	52	-38.412	93.024	-36.460	1.00	28.26	T7
	ATOM	7268	CG	PHE	A	52	-37.332	93.038	-37.504	1.00	34.93	T7
	ATOM	7269	CD1	PHE	A	52	-36.071	92.538	-37.226	1.00	31.27	T7
	ATOM	7270	CD2	PHE	A	52	-37.589	93.537	-38.777	1.00	36.38	T7
	ATOM	7271	CE1	PHE	A	52	-35.084	92.535	-38.197	1.00	24.09	T7
30	ATOM	7272	CE2	PHE	A	52	-36.609	93.536	-39.753	1.00	27.37	T7
	ATOM	7273	CZ	PHE	A	52	-35.355	93.035	-39.465	1.00	34.30	T7
	ATOM	7274	C	PHE	A	52	-39.443	93.992	-34.440	1.00	31.20	T7
	ATOM	7275	O	PHE	A	52	-39.417	93.293	-33.433	1.00	36.64	T7
	ATOM	7276	N	PHE	A	53	-40.487	94.726	-34.801	1.00	29.49	T7
35	ATOM	7277	CA	PHE	A	53	-41.752	94.716	-34.080	1.00	34.95	T7
	ATOM	7278	CB	PHE	A	53	-42.393	96.098	-34.085	1.00	28.60	T7
	ATOM	7279	CG	PHE	A	53	-43.789	96.112	-33.548	1.00	27.96	T7
	ATOM	7280	CD1	PHE	A	53	-44.034	95.857	-32.213	1.00	29.41	T7
	ATOM	7281	CD2	PHE	A	53	-44.865	96.369	-34.385	1.00	29.36	T7
40	ATOM	7282	CE1	PHE	A	53	-45.332	95.858	-31.719	1.00	24.18	T7
	ATOM	7283	CE2	PHE	A	53	-46.163	96.371	-33.897	1.00	30.77	T7
	ATOM	7284	CZ	PHE	A	53	-46.394	96.115	-32.565	1.00	28.64	T7
	ATOM	7285	C	PHE	A	53	-42.585	93.768	-34.929	1.00	29.83	T7
	ATOM	7286	O	PHE	A	53	-42.739	93.982	-36.130	1.00	33.65	T7
45	ATOM	7287	N	ILE	A	54	-43.115	92.720	-34.316	1.00	28.68	T7
	ATOM	7288	CA	ILE	A	54	-43.889	91.740	-35.060	1.00	26.85	T7
	ATOM	7289	CB	ILE	A	54	-43.188	90.376	-35.009	1.00	34.47	T7
	ATOM	7290	CG2	ILE	A	54	-43.868	89.404	-35.954	1.00	30.42	T7
	ATOM	7291	CG1	ILE	A	54	-41.717	90.550	-35.398	1.00	22.09	T7
50	ATOM	7292	CD1	ILE	A	54	-40.836	89.430	-34.959	1.00	35.47	T7
	ATOM	7293	C	ILE	A	54	-45.289	91.617	-34.502	1.00	37.02	T7
	ATOM	7294	O	ILE	A	54	-45.464	91.468	-33.302	1.00	26.47	T7
	ATOM	7295	N	TYR	A	55	-46.285	91.682	-35.378	1.00	37.53	T7
	ATOM	7296	CA	TYR	A	55	-47.671	91.590	-34.947	1.00	35.34	T7
55	ATOM	7297	CB	TYR	A	55	-48.322	92.964	-35.012	1.00	31.11	T7
	ATOM	7298	CG	TYR	A	55	-48.265	93.606	-36.369	1.00	31.47	T7
	ATOM	7299	CD1	TYR	A	55	-49.327	93.510	-37.250	1.00	28.19	T7
	ATOM	7300	CE1	TYR	A	55	-49.262	94.083	-38.512	1.00	20.44	T7
	ATOM	7301	CD2	TYR	A	55	-47.131	94.294	-36.781	1.00	31.19	T7
60	ATOM	7302	CE2	TYR	A	55	-47.054	94.870	-38.040	1.00	27.11	T7
	ATOM	7303	CZ	TYR	A	55	-48.119	94.761	-38.902	1.00	34.03	T7
	ATOM	7304	OH	TYR	A	55	-48.029	95.310	-40.158	1.00	30.93	T7
	ATOM	7305	C	TYR	A	55	-48.479	90.604	-35.767	1.00	28.85	T7
	ATOM	7306	O	TYR	A	55	-48.010	90.097	-36.771	1.00	21.60	T7
65	ATOM	7307	N	GLY	A	56	-49.697	90.322	-35.330	1.00	32.76	T7
	ATOM	7308	CA	GLY	A	56	-50.525	89.387	-36.060	1.00	28.14	T7

	ATOM	7309	C	GLY	A	56	-51.856	89.176	-35.377	1.00	27.16	T7
	ATOM	7310	O	GLY	A	56	-51.917	89.095	-34.157	1.00	33.18	T7
	ATOM	7311	N	GLN	A	57	-52.920	89.101	-36.174	1.00	31.00	T7
	ATOM	7312	CA	GLN	A	57	-54.270	88.886	-35.672	1.00	25.83	T7
5	ATOM	7313	CB	GLN	A	57	-55.087	90.186	-35.708	1.00	35.56	T7
	ATOM	7314	CG	GLN	A	57	-56.557	89.992	-35.331	1.00	30.53	T7
	ATOM	7315	CD	GLN	A	57	-57.340	91.283	-35.239	1.00	34.35	T7
	ATOM	7316	OE1	GLN	A	57	-57.149	92.072	-34.324	1.00	32.95	T7
	ATOM	7317	NE2	GLN	A	57	-58.232	91.497	-36.190	1.00	29.17	T7
10	ATOM	7318	C	GLN	A	57	-54.977	87.834	-36.509	1.00	32.73	T7
	ATOM	7319	O	GLN	A	57	-54.740	87.726	-37.706	1.00	37.93	T7
	ATOM	7320	N	VAL	A	58	-55.845	87.065	-35.861	1.00	27.95	T7
	ATOM	7321	CA	VAL	A	58	-56.620	86.014	-36.512	1.00	30.68	T7
	ATOM	7322	CB	VAL	A	58	-56.008	84.617	-36.251	1.00	28.06	T7
15	ATOM	7323	CG1	VAL	A	58	-56.961	83.531	-36.694	1.00	31.89	T7
	ATOM	7324	CG2	VAL	A	58	-54.703	84.483	-36.982	1.00	28.20	T7
	ATOM	7325	C	VAL	A	58	-58.034	86.026	-35.941	1.00	33.99	T7
	ATOM	7326	O	VAL	A	58	-58.207	86.225	-34.751	1.00	38.68	T7
	ATOM	7327	N	LEU	A	59	-59.040	85.823	-36.789	1.00	32.90	T7
20	ATOM	7328	CA	LEU	A	59	-60.430	85.786	-36.340	1.00	27.73	T7
	ATOM	7329	CB	LEU	A	59	-61.345	86.508	-37.328	1.00	30.17	T7
	ATOM	7330	CG	LEU	A	59	-62.660	87.062	-36.775	1.00	36.31	T7
	ATOM	7331	CD1	LEU	A	59	-63.599	87.347	-37.917	1.00	30.74	T7
	ATOM	7332	CD2	LEU	A	59	-63.297	86.083	-35.838	1.00	28.87	T7
25	ATOM	7333	C	LEU	A	59	-60.855	84.324	-36.243	1.00	33.85	T7
	ATOM	7334	O	LEU	A	59	-60.942	83.627	-37.253	1.00	31.20	T7
	ATOM	7335	N	TYR	A	60	-61.115	83.857	-35.029	1.00	34.20	T7
	ATOM	7336	CA	TYR	A	60	-61.529	82.476	-34.838	1.00	29.91	T7
	ATOM	7337	CB	TYR	A	60	-60.975	81.943	-33.525	1.00	37.94	T7
30	ATOM	7338	CG	TYR	A	60	-59.486	81.992	-33.485	1.00	30.23	T7
	ATOM	7339	CD1	TYR	A	60	-58.820	82.962	-32.736	1.00	30.42	T7
	ATOM	7340	CE1	TYR	A	60	-57.439	83.062	-32.770	1.00	34.41	T7
	ATOM	7341	CD2	TYR	A	60	-58.737	81.119	-34.261	1.00	21.79	T7
	ATOM	7342	CE2	TYR	A	60	-57.365	81.205	-34.311	1.00	37.32	T7
35	ATOM	7343	CZ	TYR	A	60	-56.717	82.178	-33.568	1.00	28.61	T7
	ATOM	7344	OH	TYR	A	60	-55.344	82.274	-33.641	1.00	37.95	T7
	ATOM	7345	C	TYR	A	60	-63.037	82.272	-34.861	1.00	28.86	T7
	ATOM	7346	O	TYR	A	60	-63.784	82.918	-34.130	1.00	30.95	T7
	ATOM	7347	N	THR	A	61	-63.482	81.355	-35.704	1.00	30.91	T7
40	ATOM	7348	CA	THR	A	61	-64.896	81.055	-35.810	1.00	28.84	T7
	ATOM	7349	CB	THR	A	61	-65.416	81.356	-37.219	1.00	28.81	T7
	ATOM	7350	OG1	THR	A	61	-64.615	80.673	-38.189	1.00	29.01	T7
	ATOM	7351	CG2	THR	A	61	-65.356	82.842	-37.487	1.00	36.16	T7
	ATOM	7352	C	THR	A	61	-65.093	79.585	-35.485	1.00	37.36	T7
45	ATOM	7353	O	THR	A	61	-66.124	78.997	-35.775	1.00	34.72	T7
	ATOM	7354	N	ASP	A	62	-64.073	79.005	-34.874	1.00	34.07	T7
	ATOM	7355	CA	ASP	A	62	-64.081	77.609	-34.474	1.00	27.85	T7
	ATOM	7356	CB	ASP	A	62	-62.633	77.106	-34.436	1.00	33.45	T7
	ATOM	7357	CG	ASP	A	62	-62.533	75.602	-34.344	1.00	27.88	T7
50	ATOM	7358	OD1	ASP	A	62	-61.777	75.021	-35.165	1.00	30.94	T7
	ATOM	7359	OD2	ASP	A	62	-63.197	75.012	-33.452	1.00	30.36	T7
	ATOM	7360	C	ASP	A	62	-64.719	77.551	-33.085	1.00	37.27	T7
	ATOM	7361	O	ASP	A	62	-64.599	78.496	-32.315	1.00	26.83	T7
	ATOM	7362	N	LYS	A	63	-65.401	76.463	-32.748	1.00	29.03	T7
55	ATOM	7363	CA	LYS	A	63	-66.025	76.391	-31.431	1.00	35.35	T7
	ATOM	7364	CB	LYS	A	63	-67.495	76.014	-31.569	1.00	29.31	T7
	ATOM	7365	CG	LYS	A	63	-67.728	74.671	-32.225	1.00	31.77	T7
	ATOM	7366	CD	LYS	A	63	-69.227	74.331	-32.294	1.00	32.46	T7
	ATOM	7367	CE	LYS	A	63	-70.023	75.373	-33.120	1.00	31.06	T7
60	ATOM	7368	NZ	LYS	A	63	-71.502	75.090	-33.195	1.00	36.57	T7
	ATOM	7369	C	LYS	A	63	-65.355	75.429	-30.461	1.00	31.26	T7
	ATOM	7370	O	LYS	A	63	-65.976	74.984	-29.498	1.00	32.61	T7
	ATOM	7371	N	THR	A	64	-64.086	75.120	-30.685	1.00	35.97	T7
	ATOM	7372	CA	THR	A	64	-63.420	74.185	-29.796	1.00	32.33	T7
65	ATOM	7373	CB	THR	A	64	-62.324	73.371	-30.546	1.00	28.61	T7
	ATOM	7374	OG1	THR	A	64	-61.446	74.253	-31.251	1.00	31.36	T7

	ATOM	7375	CG2	THR	A	64	-62.972	72.403	-31.530	1.00	32.10	T7
	ATOM	7376	C	THR	A	64	-62.839	74.757	-28.506	1.00	36.01	T7
	ATOM	7377	O	THR	A	64	-61.644	74.663	-28.263	1.00	30.40	T7
	ATOM	7378	N	TYR	A	65	-63.705	75.337	-27.684	1.00	29.48	T7
5	ATOM	7379	CA	TYR	A	65	-63.341	75.898	-26.376	1.00	28.29	T7
	ATOM	7380	CB	TYR	A	65	-63.251	74.762	-25.348	1.00	24.60	T7
	ATOM	7381	CG	TYR	A	65	-61.847	74.336	-24.988	1.00	36.52	T7
	ATOM	7382	CD1	TYR	A	65	-61.174	74.910	-23.906	1.00	27.88	T7
	ATOM	7383	CE1	TYR	A	65	-59.864	74.522	-23.576	1.00	30.50	T7
10	ATOM	7384	CD2	TYR	A	65	-61.181	73.366	-25.734	1.00	29.19	T7
	ATOM	7385	CE2	TYR	A	65	-59.872	72.972	-25.417	1.00	22.79	T7
	ATOM	7386	CZ	TYR	A	65	-59.223	73.550	-24.342	1.00	31.04	T7
	ATOM	7387	OH	TYR	A	65	-57.941	73.147	-24.043	1.00	34.37	T7
	ATOM	7388	C	TYR	A	65	-62.100	76.794	-26.235	1.00	36.58	T7
15	ATOM	7389	O	TYR	A	65	-62.030	77.600	-25.303	1.00	31.90	T7
	ATOM	7390	N	ALA	A	66	-61.125	76.661	-27.128	1.00	32.94	T7
	ATOM	7391	CA	ALA	A	66	-59.924	77.475	-27.033	1.00	25.53	T7
	ATOM	7392	CB	ALA	A	66	-59.046	76.966	-25.904	1.00	33.91	T7
	ATOM	7393	C	ALA	A	66	-59.143	77.496	-28.340	1.00	32.43	T7
20	ATOM	7394	O	ALA	A	66	-58.677	76.467	-28.817	1.00	32.77	T7
	ATOM	7395	N	MET	A	67	-59.008	78.678	-28.924	1.00	27.32	T7
	ATOM	7396	CA	MET	A	67	-58.272	78.828	-30.163	1.00	29.70	T7
	ATOM	7397	CB	MET	A	67	-59.192	79.341	-31.267	1.00	30.99	T7
	ATOM	7398	CG	MET	A	67	-60.230	78.323	-31.723	1.00	24.78	T7
25	ATOM	7399	SD	MET	A	67	-59.457	76.822	-32.363	1.00	37.32	T7
	ATOM	7400	CE	MET	A	67	-59.010	77.346	-34.017	1.00	30.54	T7
	ATOM	7401	C	MET	A	67	-57.137	79.806	-29.942	1.00	27.89	T7
	ATOM	7402	O	MET	A	67	-57.096	80.489	-28.925	1.00	32.47	T7
	ATOM	7403	N	GLY	A	68	-56.211	79.866	-30.893	1.00	27.91	T7
30	ATOM	7404	CA	GLY	A	68	-55.083	80.774	-30.772	1.00	24.32	T7
	ATOM	7405	C	GLY	A	68	-54.025	80.524	-31.826	1.00	29.89	T7
	ATOM	7406	O	GLY	A	68	-54.089	79.531	-32.547	1.00	29.42	T7
	ATOM	7407	N	HIS	A	69	-53.061	81.430	-31.936	1.00	33.42	T7
	ATOM	7408	CA	HIS	A	69	-51.989	81.269	-32.907	1.00	39.72	T7
35	ATOM	7409	CB	HIS	A	69	-52.231	82.128	-34.152	1.00	37.37	T7
	ATOM	7410	CG	HIS	A	69	-52.458	83.580	-33.871	1.00	26.47	T7
	ATOM	7411	CD2	HIS	A	69	-51.649	84.653	-34.024	1.00	37.58	T7
	ATOM	7412	ND1	HIS	A	69	-53.659	84.073	-33.411	1.00	28.35	T7
	ATOM	7413	CE1	HIS	A	69	-53.583	85.386	-33.298	1.00	32.16	T7
40	ATOM	7414	NE2	HIS	A	69	-52.374	85.763	-33.664	1.00	27.21	T7
	ATOM	7415	C	HIS	A	69	-50.615	81.574	-32.328	1.00	30.30	T7
	ATOM	7416	O	HIS	A	69	-50.494	82.124	-31.237	1.00	30.50	T7
	ATOM	7417	N	LEU	A	70	-49.582	81.200	-33.072	1.00	34.64	T7
	ATOM	7418	CA	LEU	A	70	-48.203	81.401	-32.653	1.00	32.49	T7
45	ATOM	7419	CB	LEU	A	70	-47.502	80.058	-32.492	1.00	35.68	T7
	ATOM	7420	CG	LEU	A	70	-48.269	78.908	-31.864	1.00	32.27	T7
	ATOM	7421	CD1	LEU	A	70	-47.453	77.656	-32.001	1.00	31.25	T7
	ATOM	7422	CD2	LEU	A	70	-48.560	79.203	-30.417	1.00	32.86	T7
	ATOM	7423	C	LEU	A	70	-47.437	82.175	-33.703	1.00	26.26	T7
50	ATOM	7424	O	LEU	A	70	-47.673	82.006	-34.891	1.00	26.73	T7
	ATOM	7425	N	ILE	A	71	-46.522	83.027	-33.267	1.00	30.34	T7
	ATOM	7426	CA	ILE	A	71	-45.687	83.761	-34.199	1.00	31.36	T7
	ATOM	7427	CB	ILE	A	71	-45.650	85.257	-33.876	1.00	31.55	T7
	ATOM	7428	CG2	ILE	A	71	-44.539	85.927	-34.644	1.00	34.44	T7
55	ATOM	7429	CG1	ILE	A	71	-46.988	85.893	-34.244	1.00	32.73	T7
	ATOM	7430	CD1	ILE	A	71	-47.069	87.365	-33.937	1.00	27.98	T7
	ATOM	7431	C	ILE	A	71	-44.325	83.128	-33.974	1.00	27.75	T7
	ATOM	7432	O	ILE	A	71	-43.710	83.310	-32.929	1.00	32.14	T7
	ATOM	7433	N	GLN	A	72	-43.851	82.368	-34.952	1.00	33.38	T7
60	ATOM	7434	CA	GLN	A	72	-42.581	81.674	-34.790	1.00	30.56	T7
	ATOM	7435	CB	GLN	A	72	-42.792	80.192	-35.049	1.00	35.89	T7
	ATOM	7436	CG	GLN	A	72	-44.015	79.666	-34.350	1.00	28.69	T7
	ATOM	7437	CD	GLN	A	72	-44.118	78.171	-34.442	1.00	32.84	T7
	ATOM	7438	OE1	GLN	A	72	-44.084	77.603	-35.534	1.00	37.74	T7
65	ATOM	7439	NE2	GLN	A	72	-44.245	77.515	-33.295	1.00	27.32	T7
	ATOM	7440	C	GLN	A	72	-41.418	82.172	-35.626	1.00	28.34	T7

	ATOM	7441	O	GLN	A	72	-41.601	82.799	-36.669	1.00	25.20	T7
	ATOM	7442	N	ARG	A	73	-40.218	81.861	-35.149	1.00	28.62	T7
	ATOM	7443	CA	ARG	A	73	-38.980	82.259	-35.797	1.00	28.52	T7
	ATOM	7444	CB	ARG	A	73	-38.213	83.191	-34.869	1.00	32.20	T7
5	ATOM	7445	CG	ARG	A	73	-36.904	83.646	-35.422	1.00	36.48	T7
	ATOM	7446	CD	ARG	A	73	-35.959	84.019	-34.314	1.00	24.34	T7
	ATOM	7447	NE	ARG	A	73	-34.671	84.456	-34.837	1.00	31.85	T7
	ATOM	7448	CZ	ARG	A	73	-33.578	84.597	-34.101	1.00	34.42	T7
	ATOM	7449	NH1	ARG	A	73	-33.607	84.333	-32.804	1.00	28.25	T7
10	ATOM	7450	NH2	ARG	A	73	-32.456	85.009	-34.664	1.00	33.67	T7
	ATOM	7451	C	ARG	A	73	-38.111	81.040	-36.122	1.00	27.22	T7
	ATOM	7452	O	ARG	A	73	-37.916	80.169	-35.275	1.00	29.04	T7
	ATOM	7453	N	LYS	A	74	-37.600	80.978	-37.351	1.00	27.59	T7
	ATOM	7454	CA	LYS	A	74	-36.724	79.882	-37.774	1.00	29.93	T7
15	ATOM	7455	CB	LYS	A	74	-37.130	79.351	-39.143	1.00	37.07	T7
	ATOM	7456	CG	LYS	A	74	-38.463	78.634	-39.166	1.00	33.97	T7
	ATOM	7457	CD	LYS	A	74	-38.866	78.196	-40.594	1.00	27.87	T7
	ATOM	7458	CE	LYS	A	74	-40.233	77.493	-40.603	1.00	38.08	T7
	ATOM	7459	NZ	LYS	A	74	-40.638	77.019	-41.964	1.00	35.64	T7
20	ATOM	7460	C	LYS	A	74	-35.304	80.418	-37.863	1.00	27.93	T7
	ATOM	7461	O	LYS	A	74	-34.940	81.078	-38.843	1.00	33.00	T7
	ATOM	7462	N	LYS	A	75	-34.505	80.138	-36.839	1.00	29.00	T7
	ATOM	7463	CA	LYS	A	75	-33.126	80.602	-36.787	1.00	36.16	T7
	ATOM	7464	CB	LYS	A	75	-32.462	80.132	-35.490	1.00	34.22	T7
25	ATOM	7465	CG	LYS	A	75	-33.088	80.625	-34.203	1.00	21.63	T7
	ATOM	7466	CD	LYS	A	75	-32.318	80.072	-33.015	1.00	32.04	T7
	ATOM	7467	CE	LYS	A	75	-32.921	80.544	-31.699	1.00	32.31	T7
	ATOM	7468	NZ	LYS	A	75	-32.274	79.962	-30.469	1.00	30.05	T7
	ATOM	7469	C	LYS	A	75	-32.313	80.074	-37.960	1.00	27.85	T7
30	ATOM	7470	O	LYS	A	75	-32.419	78.898	-38.304	1.00	26.60	T7
	ATOM	7471	N	VAL	A	76	-31.493	80.933	-38.562	1.00	30.99	T7
	ATOM	7472	CA	VAL	A	76	-30.644	80.514	-39.677	1.00	30.50	T7
	ATOM	7473	CB	VAL	A	76	-30.059	81.692	-40.452	1.00	29.05	T7
	ATOM	7474	CG1	VAL	A	76	-29.564	81.215	-41.791	1.00	33.08	T7
35	ATOM	7475	CG2	VAL	A	76	-31.072	82.765	-40.606	1.00	32.72	T7
	ATOM	7476	C	VAL	A	76	-29.455	79.790	-39.088	1.00	29.98	T7
	ATOM	7477	O	VAL	A	76	-29.004	78.772	-39.607	1.00	28.82	T7
	ATOM	7478	N	HIS	A	77	-28.948	80.351	-37.999	1.00	31.20	T7
	ATOM	7479	CA	HIS	A	77	-27.807	79.798	-37.304	1.00	30.05	T7
40	ATOM	7480	CB	HIS	A	77	-26.828	80.913	-36.969	1.00	37.65	T7
	ATOM	7481	CG	HIS	A	77	-26.348	81.663	-38.170	1.00	34.19	T7
	ATOM	7482	CD2	HIS	A	77	-25.800	82.895	-38.294	1.00	30.81	T7
	ATOM	7483	ND1	HIS	A	77	-26.348	81.112	-39.435	1.00	34.05	T7
	ATOM	7484	CE1	HIS	A	77	-25.817	81.969	-40.286	1.00	30.85	T7
45	ATOM	7485	NE2	HIS	A	77	-25.476	83.058	-39.618	1.00	37.36	T7
	ATOM	7486	C	HIS	A	77	-28.278	79.105	-36.041	1.00	31.82	T7
	ATOM	7487	O	HIS	A	77	-29.235	79.547	-35.417	1.00	26.59	T7
	ATOM	7488	N	VAL	A	78	-27.599	78.034	-35.643	1.00	31.20	T7
	ATOM	7489	CA	VAL	A	78	-28.048	77.308	-34.477	1.00	26.02	T7
50	ATOM	7490	CB	VAL	A	78	-28.656	75.964	-34.922	1.00	28.39	T7
	ATOM	7491	CG1	VAL	A	78	-29.269	75.244	-33.749	1.00	29.04	T7
	ATOM	7492	CG2	VAL	A	78	-29.736	76.218	-35.949	1.00	26.91	T7
	ATOM	7493	C	VAL	A	78	-27.105	77.101	-33.282	1.00	29.50	T7
	ATOM	7494	O	VAL	A	78	-27.376	77.633	-32.202	1.00	28.43	T7
55	ATOM	7495	N	PHE	A	79	-26.022	76.346	-33.423	1.00	28.23	T7
	ATOM	7496	CA	PHE	A	79	-25.118	76.118	-32.270	1.00	32.20	T7
	ATOM	7497	CB	PHE	A	79	-24.840	77.407	-31.476	1.00	25.88	T7
	ATOM	7498	CG	PHE	A	79	-24.366	78.552	-32.307	1.00	28.22	T7
	ATOM	7499	CD1	PHE	A	79	-25.243	79.567	-32.678	1.00	35.25	T7
60	ATOM	7500	CD2	PHE	A	79	-23.044	78.618	-32.725	1.00	27.86	T7
	ATOM	7501	CE1	PHE	A	79	-24.812	80.634	-33.455	1.00	36.47	T7
	ATOM	7502	CE2	PHE	A	79	-22.599	79.678	-33.504	1.00	24.70	T7
	ATOM	7503	CZ	PHE	A	79	-23.486	80.692	-33.871	1.00	25.73	T7
	ATOM	7504	C	PHE	A	79	-25.626	75.093	-31.242	1.00	25.00	T7
65	ATOM	7505	O	PHE	A	79	-26.714	75.242	-30.674	1.00	29.90	T7
	ATOM	7506	N	GLY	A	80	-24.807	74.075	-30.989	1.00	29.55	T7

	ATOM	7507	CA	GLY	A	80	-25.138	73.049	-30.015	1.00	28.57	T7
	ATOM	7508	C	GLY	A	80	-26.537	72.466	-30.069	1.00	31.89	T7
	ATOM	7509	O	GLY	A	80	-27.018	72.052	-31.130	1.00	32.27	T7
	ATOM	7510	N	ASP	A	81	-27.191	72.437	-28.908	1.00	28.78	T7
5	ATOM	7511	CA	ASP	A	81	-28.536	71.879	-28.791	1.00	28.66	T7
	ATOM	7512	CB	ASP	A	81	-28.635	71.031	-27.516	1.00	26.09	T7
	ATOM	7513	CG	ASP	A	81	-28.491	71.855	-26.249	1.00	27.39	T7
	ATOM	7514	OD1	ASP	A	81	-28.009	73.008	-26.331	1.00	28.83	T7
	ATOM	7515	OD2	ASP	A	81	-28.848	71.349	-25.162	1.00	25.86	T7
10	ATOM	7516	C	ASP	A	81	-29.659	72.917	-28.816	1.00	33.95	T7
	ATOM	7517	O	ASP	A	81	-30.712	72.718	-28.206	1.00	26.27	T7
	ATOM	7518	N	GLU	A	82	-29.434	74.029	-29.510	1.00	26.51	T7
	ATOM	7519	CA	GLU	A	82	-30.465	75.055	-29.616	1.00	28.09	T7
	ATOM	7520	CB	GLU	A	82	-29.936	76.307	-30.304	1.00	26.66	T7
15	ATOM	7521	CG	GLU	A	82	-29.132	77.245	-29.483	1.00	30.25	T7
	ATOM	7522	CD	GLU	A	82	-29.264	78.649	-30.023	1.00	28.33	T7
	ATOM	7523	OE1	GLU	A	82	-29.199	78.809	-31.259	1.00	31.41	T7
	ATOM	7524	OE2	GLU	A	82	-29.441	79.595	-29.223	1.00	29.35	T7
	ATOM	7525	C	GLU	A	82	-31.552	74.495	-30.517	1.00	31.05	T7
20	ATOM	7526	O	GLU	A	82	-31.260	73.684	-31.395	1.00	33.23	T7
	ATOM	7527	N	LEU	A	83	-32.796	74.909	-30.309	1.00	25.56	T7
	ATOM	7528	CA	LEU	A	83	-33.864	74.463	-31.195	1.00	34.37	T7
	ATOM	7529	CB	LEU	A	83	-35.206	74.370	-30.472	1.00	29.92	T7
	ATOM	7530	CG	LEU	A	83	-35.359	73.268	-29.431	1.00	38.98	T7
25	ATOM	7531	CD1	LEU	A	83	-34.396	73.508	-28.281	1.00	30.48	T7
	ATOM	7532	CD2	LEU	A	83	-36.775	73.266	-28.925	1.00	34.91	T7
	ATOM	7533	C	LEU	A	83	-33.900	75.591	-32.200	1.00	33.69	T7
	ATOM	7534	O	LEU	A	83	-33.913	76.761	-31.816	1.00	35.47	T7
	ATOM	7535	N	SER	A	84	-33.897	75.254	-33.480	1.00	31.58	T7
30	ATOM	7536	CA	SER	A	84	-33.898	76.287	-34.506	1.00	29.48	T7
	ATOM	7537	CB	SER	A	84	-33.359	75.728	-35.828	1.00	26.83	T7
	ATOM	7538	OG	SER	A	84	-33.965	74.491	-36.133	1.00	27.79	T7
	ATOM	7539	C	SER	A	84	-35.257	76.930	-34.728	1.00	32.93	T7
	ATOM	7540	O	SER	A	84	-35.381	77.870	-35.518	1.00	25.28	T7
35	ATOM	7541	N	LEU	A	85	-36.272	76.436	-34.027	1.00	26.11	T7
	ATOM	7542	CA	LEU	A	85	-37.612	76.985	-34.163	1.00	33.20	T7
	ATOM	7543	CB	LEU	A	85	-38.583	75.896	-34.627	1.00	27.45	T7
	ATOM	7544	CG	LEU	A	85	-39.960	76.267	-35.204	1.00	24.21	T7
	ATOM	7545	CD1	LEU	A	85	-40.871	76.802	-34.122	1.00	29.88	T7
40	ATOM	7546	CD2	LEU	A	85	-39.792	77.288	-36.322	1.00	25.19	T7
	ATOM	7547	C	LEU	A	85	-38.038	77.534	-32.822	1.00	29.62	T7
	ATOM	7548	O	LEU	A	85	-38.309	76.787	-31.894	1.00	26.70	T7
	ATOM	7549	N	VAL	A	86	-38.071	78.851	-32.712	1.00	27.37	T7
	ATOM	7550	CA	VAL	A	86	-38.466	79.497	-31.467	1.00	33.74	T7
45	ATOM	7551	CB	VAL	A	86	-37.480	80.616	-31.053	1.00	24.40	T7
	ATOM	7552	CG1	VAL	A	86	-37.997	81.325	-29.817	1.00	40.49	T7
	ATOM	7553	CG2	VAL	A	86	-36.111	80.034	-30.786	1.00	33.48	T7
	ATOM	7554	C	VAL	A	86	-39.813	80.141	-31.675	1.00	27.29	T7
	ATOM	7555	O	VAL	A	86	-40.067	80.732	-32.719	1.00	30.95	T7
50	ATOM	7556	N	THR	A	87	-40.686	80.024	-30.687	1.00	25.16	T7
	ATOM	7557	CA	THR	A	87	-41.983	80.649	-30.814	1.00	29.99	T7
	ATOM	7558	CB	THR	A	87	-43.127	79.657	-30.501	1.00	31.94	T7
	ATOM	7559	OG1	THR	A	87	-43.863	80.116	-29.373	1.00	34.22	T7
	ATOM	7560	CG2	THR	A	87	-42.575	78.270	-30.229	1.00	33.56	T7
55	ATOM	7561	C	THR	A	87	-42.035	81.867	-29.892	1.00	28.80	T7
	ATOM	7562	O	THR	A	87	-41.895	81.758	-28.680	1.00	31.02	T7
	ATOM	7563	N	LEU	A	88	-42.198	83.036	-30.501	1.00	32.09	T7
	ATOM	7564	CA	LEU	A	88	-42.281	84.302	-29.790	1.00	36.00	T7
	ATOM	7565	CB	LEU	A	88	-41.651	85.409	-30.637	1.00	35.69	T7
60	ATOM	7566	CG	LEU	A	88	-40.225	85.379	-31.198	1.00	32.01	T7
	ATOM	7567	CD1	LEU	A	88	-39.758	83.983	-31.435	1.00	34.87	T7
	ATOM	7568	CD2	LEU	A	88	-40.192	86.151	-32.498	1.00	25.35	T7
	ATOM	7569	C	LEU	A	88	-43.764	84.635	-29.627	1.00	32.08	T7
	ATOM	7570	O	LEU	A	88	-44.585	84.314	-30.482	1.00	30.71	T7
65	ATOM	7571	N	PHE	A	89	-44.143	85.256	-28.530	1.00	36.07	T7
	ATOM	7572	CA	PHE	A	89	-45.543	85.668	-28.402	1.00	28.92	T7

	ATOM	7573	CB	PHE	A	89	-45.873	86.613	-29.562	1.00	30.86	T7
	ATOM	7574	CG	PHE	A	89	-44.779	87.599	-29.833	1.00	23.36	T7
	ATOM	7575	CD1	PHE	A	89	-44.503	88.017	-31.126	1.00	36.06	T7
	ATOM	7576	CD2	PHE	A	89	-43.942	88.040	-28.788	1.00	33.33	T7
5	ATOM	7577	CE1	PHE	A	89	-43.398	88.852	-31.380	1.00	29.58	T7
	ATOM	7578	CE2	PHE	A	89	-42.844	88.868	-29.027	1.00	29.55	T7
	ATOM	7579	CZ	PHE	A	89	-42.568	89.273	-30.322	1.00	30.94	T7
	ATOM	7580	C	PHE	A	89	-46.639	84.610	-28.254	1.00	33.41	T7
	ATOM	7581	O	PHE	A	89	-46.784	84.030	-27.178	1.00	30.00	T7
10	ATOM	7582	N	ARG	A	90	-47.442	84.371	-29.286	1.00	31.01	T7
	ATOM	7583	CA	ARG	A	90	-48.528	83.388	-29.132	1.00	31.58	T7
	ATOM	7584	CB	ARG	A	90	-47.965	82.062	-28.619	1.00	29.78	T7
	ATOM	7585	CG	ARG	A	90	-48.953	81.230	-27.836	1.00	28.84	T7
	ATOM	7586	CD	ARG	A	90	-48.244	80.153	-27.039	1.00	31.62	T7
15	ATOM	7587	NE	ARG	A	90	-49.170	79.513	-26.110	1.00	27.17	T7
	ATOM	7588	CZ	ARG	A	90	-48.846	79.100	-24.886	1.00	29.83	T7
	ATOM	7589	NH1	ARG	A	90	-47.607	79.252	-24.433	1.00	35.04	T7
	ATOM	7590	NH2	ARG	A	90	-49.774	78.556	-24.106	1.00	29.59	T7
	ATOM	7591	C	ARG	A	90	-49.662	83.825	-28.183	1.00	33.87	T7
20	ATOM	7592	O	ARG	A	90	-49.415	84.194	-27.040	1.00	34.78	T7
	ATOM	7593	N	CYS	A	91	-50.903	83.746	-28.660	1.00	30.02	T7
	ATOM	7594	CA	CYS	A	91	-52.072	84.119	-27.862	1.00	27.25	T7
	ATOM	7595	CB	CYS	A	91	-52.655	85.431	-28.375	1.00	36.20	T7
	ATOM	7596	SG	CYS	A	91	-53.084	85.389	-30.101	1.00	30.17	T7
25	ATOM	7597	C	CYS	A	91	-53.156	83.029	-27.857	1.00	28.07	T7
	ATOM	7598	O	CYS	A	91	-53.117	82.105	-28.665	1.00	27.87	T7
	ATOM	7599	N	ILE	A	92	-54.120	83.146	-26.941	1.00	30.24	T7
	ATOM	7600	CA	ILE	A	92	-55.203	82.166	-26.796	1.00	34.47	T7
	ATOM	7601	CB	ILE	A	92	-54.947	81.237	-25.599	1.00	26.67	T7
30	ATOM	7602	CG2	ILE	A	92	-56.026	80.178	-25.517	1.00	32.70	T7
	ATOM	7603	CG1	ILE	A	92	-53.582	80.566	-25.747	1.00	30.57	T7
	ATOM	7604	CD1	ILE	A	92	-53.104	79.863	-24.485	1.00	26.64	T7
	ATOM	7605	C	ILE	A	92	-56.527	82.859	-26.534	1.00	32.34	T7
	ATOM	7606	O	ILE	A	92	-56.550	83.955	-26.014	1.00	28.09	T7
35	ATOM	7607	N	GLN	A	93	-57.631	82.217	-26.889	1.00	35.89	T7
	ATOM	7608	CA	GLN	A	93	-58.955	82.785	-26.658	1.00	32.56	T7
	ATOM	7609	CB	GLN	A	93	-59.371	83.690	-27.817	1.00	40.16	T7
	ATOM	7610	CG	GLN	A	93	-58.892	85.125	-27.698	1.00	32.25	T7
	ATOM	7611	CD	GLN	A	93	-59.946	86.060	-27.138	1.00	28.94	T7
40	ATOM	7612	OE1	GLN	A	93	-60.228	86.066	-25.941	1.00	30.79	T7
	ATOM	7613	NE2	GLN	A	93	-60.548	86.854	-28.016	1.00	34.20	T7
	ATOM	7614	C	GLN	A	93	-59.995	81.689	-26.484	1.00	30.59	T7
	ATOM	7615	O	GLN	A	93	-60.191	80.867	-27.378	1.00	26.72	T7
	ATOM	7616	N	ASN	A	94	-60.646	81.667	-25.324	1.00	27.78	T7
45	ATOM	7617	CA	ASN	A	94	-61.679	80.677	-25.073	1.00	35.85	T7
	ATOM	7618	CB	ASN	A	94	-62.262	80.852	-23.670	1.00	28.88	T7
	ATOM	7619	CG	ASN	A	94	-61.402	80.227	-22.598	1.00	33.36	T7
	ATOM	7620	OD1	ASN	A	94	-61.086	79.053	-22.672	1.00	28.51	T7
	ATOM	7621	ND2	ASN	A	94	-61.032	81.001	-21.588	1.00	25.09	T7
50	ATOM	7622	C	ASN	A	94	-62.755	80.924	-26.124	1.00	23.76	T7
	ATOM	7623	O	ASN	A	94	-62.984	82.067	-26.530	1.00	34.36	T7
	ATOM	7624	N	MET	A	95	-63.405	79.859	-26.578	1.00	35.51	T7
	ATOM	7625	CA	MET	A	95	-64.452	79.990	-27.579	1.00	29.96	T7
	ATOM	7626	CB	MET	A	95	-64.140	79.110	-28.794	1.00	26.32	T7
55	ATOM	7627	CG	MET	A	95	-62.859	79.454	-29.525	1.00	35.54	T7
	ATOM	7628	SD	MET	A	95	-62.779	81.192	-29.983	1.00	24.99	T7
	ATOM	7629	CE	MET	A	95	-63.897	81.303	-31.335	1.00	34.72	T7
	ATOM	7630	C	MET	A	95	-65.803	79.580	-27.006	1.00	33.86	T7
	ATOM	7631	O	MET	A	95	-65.873	78.765	-26.085	1.00	34.55	T7
60	ATOM	7632	N	PRO	A	96	-66.895	80.159	-27.534	1.00	28.07	T7
	ATOM	7633	CD	PRO	A	96	-66.926	81.251	-28.514	1.00	27.65	T7
	ATOM	7634	CA	PRO	A	96	-68.255	79.847	-27.088	1.00	35.61	T7
	ATOM	7635	CB	PRO	A	96	-69.100	80.976	-27.671	1.00	21.50	T7
	ATOM	7636	CG	PRO	A	96	-68.110	82.032	-28.030	1.00	29.62	T7
65	ATOM	7637	C	PRO	A	96	-68.608	78.529	-27.756	1.00	38.58	T7
	ATOM	7638	O	PRO	A	96	-67.750	77.874	-28.366	1.00	31.29	T7

	ATOM	7639	N	GLU	A	97	-69.873	78.150	-27.679	1.00	32.55	T7
	ATOM	7640	CA	GLU	A	97	-70.283	76.902	-28.287	1.00	25.91	T7
	ATOM	7641	CB	GLU	A	97	-70.942	76.022	-27.240	1.00	27.77	T7
	ATOM	7642	CG	GLU	A	97	-71.056	74.586	-27.665	1.00	23.75	T7
5	ATOM	7643	CD	GLU	A	97	-70.674	73.649	-26.535	1.00	31.85	T7
	ATOM	7644	OE1	GLU	A	97	-71.340	73.700	-25.469	1.00	31.89	T7
	ATOM	7645	OE2	GLU	A	97	-69.699	72.868	-26.707	1.00	30.25	T7
	ATOM	7646	C	GLU	A	97	-71.249	77.172	-29.418	1.00	30.67	T7
	ATOM	7647	O	GLU	A	97	-71.400	76.364	-30.339	1.00	28.89	T7
10	ATOM	7648	N	THR	A	98	-71.881	78.334	-29.359	1.00	29.97	T7
	ATOM	7649	CA	THR	A	98	-72.859	78.698	-30.358	1.00	28.87	T7
	ATOM	7650	CB	THR	A	98	-73.987	79.471	-29.730	1.00	30.88	T7
	ATOM	7651	OG1	THR	A	98	-73.455	80.699	-29.219	1.00	21.20	T7
	ATOM	7652	CG2	THR	A	98	-74.616	78.659	-28.599	1.00	30.04	T7
15	ATOM	7653	C	THR	A	98	-72.339	79.514	-31.527	1.00	26.70	T7
	ATOM	7654	O	THR	A	98	-72.238	78.991	-32.637	1.00	26.00	T7
	ATOM	7655	N	LEU	A	99	-72.007	80.780	-31.311	1.00	36.25	T7
	ATOM	7656	CA	LEU	A	99	-71.570	81.580	-32.441	1.00	28.78	T7
	ATOM	7657	CB	LEU	A	99	-72.534	82.746	-32.634	1.00	33.22	T7
20	ATOM	7658	CG	LEU	A	99	-73.955	82.273	-32.965	1.00	33.09	T7
	ATOM	7659	CD1	LEU	A	99	-74.924	83.442	-32.986	1.00	33.28	T7
	ATOM	7660	CD2	LEU	A	99	-73.941	81.571	-34.314	1.00	34.50	T7
	ATOM	7661	C	LEU	A	99	-70.144	82.078	-32.358	1.00	26.84	T7
	ATOM	7662	O	LEU	A	99	-69.902	83.275	-32.153	1.00	26.42	T7
25	ATOM	7663	N	PRO	A	100	-69.172	81.166	-32.544	1.00	40.65	T7
	ATOM	7664	CD	PRO	A	100	-69.365	79.752	-32.907	1.00	37.11	T7
	ATOM	7665	CA	PRO	A	100	-67.744	81.482	-32.493	1.00	29.65	T7
	ATOM	7666	CB	PRO	A	100	-67.106	80.231	-33.080	1.00	29.02	T7
	ATOM	7667	CG	PRO	A	100	-68.015	79.163	-32.605	1.00	28.71	T7
30	ATOM	7668	C	PRO	A	100	-67.373	82.740	-33.270	1.00	29.95	T7
	ATOM	7669	O	PRO	A	100	-67.624	82.841	-34.470	1.00	30.63	T7
	ATOM	7670	N	ASN	A	101	-66.772	83.691	-32.570	1.00	32.36	T7
	ATOM	7671	CA	ASN	A	101	-66.349	84.943	-33.166	1.00	26.74	T7
	ATOM	7672	CB	ASN	A	101	-67.536	85.856	-33.374	1.00	31.37	T7
35	ATOM	7673	CG	ASN	A	101	-68.200	85.631	-34.677	1.00	33.74	T7
	ATOM	7674	OD1	ASN	A	101	-67.634	85.938	-35.721	1.00	31.16	T7
	ATOM	7675	ND2	ASN	A	101	-69.412	85.081	-34.644	1.00	28.47	T7
	ATOM	7676	C	ASN	A	101	-65.398	85.640	-32.232	1.00	30.48	T7
	ATOM	7677	O	ASN	A	101	-65.824	86.515	-31.475	1.00	32.39	T7
40	ATOM	7678	N	ASN	A	102	-64.115	85.284	-32.260	1.00	30.26	T7
	ATOM	7679	CA	ASN	A	102	-63.229	85.967	-31.352	1.00	32.18	T7
	ATOM	7680	CB	ASN	A	102	-62.740	85.009	-30.286	1.00	29.82	T7
	ATOM	7681	CG	ASN	A	102	-63.739	84.890	-29.148	1.00	26.14	T7
	ATOM	7682	OD1	ASN	A	102	-64.269	85.896	-28.676	1.00	24.59	T7
45	ATOM	7683	ND2	ASN	A	102	-64.006	83.666	-28.707	1.00	22.15	T7
	ATOM	7684	C	ASN	A	102	-62.106	86.856	-31.841	1.00	29.33	T7
	ATOM	7685	O	ASN	A	102	-61.983	87.970	-31.332	1.00	27.30	T7
	ATOM	7686	N	SER	A	103	-61.291	86.452	-32.799	1.00	30.76	T7
	ATOM	7687	CA	SER	A	103	-60.240	87.396	-33.203	1.00	36.12	T7
50	ATOM	7688	CB	SER	A	103	-60.866	88.689	-33.762	1.00	29.97	T7
	ATOM	7689	OG	SER	A	103	-60.268	89.852	-33.210	1.00	26.98	T7
	ATOM	7690	C	SER	A	103	-59.302	87.745	-32.027	1.00	38.41	T7
	ATOM	7691	O	SER	A	103	-59.731	88.116	-30.936	1.00	28.64	T7
	ATOM	7692	N	CYS	A	104	-58.006	87.635	-32.259	1.00	29.03	T7
55	ATOM	7693	CA	CYS	A	104	-57.063	87.911	-31.208	1.00	30.84	T7
	ATOM	7694	CB	CYS	A	104	-56.760	86.618	-30.475	1.00	28.77	T7
	ATOM	7695	SG	CYS	A	104	-55.860	86.820	-28.977	1.00	28.38	T7
	ATOM	7696	C	CYS	A	104	-55.797	88.496	-31.801	1.00	36.75	T7
	ATOM	7697	O	CYS	A	104	-55.215	87.936	-32.724	1.00	29.83	T7
60	ATOM	7698	N	TYR	A	105	-55.388	89.638	-31.267	1.00	24.30	T7
	ATOM	7699	CA	TYR	A	105	-54.192	90.327	-31.719	1.00	29.74	T7
	ATOM	7700	CB	TYR	A	105	-54.493	91.811	-31.902	1.00	26.47	T7
	ATOM	7701	CG	TYR	A	105	-53.299	92.662	-32.276	1.00	31.54	T7
	ATOM	7702	CD1	TYR	A	105	-53.012	92.951	-33.606	1.00	26.98	T7
65	ATOM	7703	CE1	TYR	A	105	-51.931	93.749	-33.951	1.00	30.66	T7
	ATOM	7704	CD2	TYR	A	105	-52.467	93.192	-31.300	1.00	27.61	T7

	ATOM	7705	CE2	TYR	A	105	-51.378	93.989	-31.639	1.00	30.88	T7
	ATOM	7706	CZ	TYR	A	105	-51.121	94.264	-32.964	1.00	36.49	T7
	ATOM	7707	OH	TYR	A	105	-50.064	95.067	-33.301	1.00	29.44	T7
	ATOM	7708	C	TYR	A	105	-53.098	90.174	-30.683	1.00	26.54	T7
5	ATOM	7709	O	TYR	A	105	-53.369	90.114	-29.490	1.00	32.15	T7
	ATOM	7710	N	SER	A	106	-51.856	90.112	-31.137	1.00	29.69	T7
	ATOM	7711	CA	SER	A	106	-50.726	90.001	-30.227	1.00	30.66	T7
	ATOM	7712	CB	SER	A	106	-50.565	88.565	-29.744	1.00	30.95	T7
	ATOM	7713	OG	SER	A	106	-49.625	88.492	-28.695	1.00	33.48	T7
10	ATOM	7714	C	SER	A	106	-49.504	90.449	-30.998	1.00	32.17	T7
	ATOM	7715	O	SER	A	106	-49.414	90.207	-32.198	1.00	27.08	T7
	ATOM	7716	N	ALA	A	107	-48.583	91.125	-30.318	1.00	28.14	T7
	ATOM	7717	CA	ALA	A	107	-47.369	91.626	-30.954	1.00	28.45	T7
	ATOM	7718	CB	ALA	A	107	-47.649	92.946	-31.643	1.00	36.55	T7
15	ATOM	7719	C	ALA	A	107	-46.253	91.802	-29.947	1.00	27.73	T7
	ATOM	7720	O	ALA	A	107	-46.486	91.806	-28.746	1.00	32.40	T7
	ATOM	7721	N	GLY	A	108	-45.035	91.945	-30.447	1.00	30.13	T7
	ATOM	7722	CA	GLY	A	108	-43.886	92.125	-29.580	1.00	33.86	T7
	ATOM	7723	C	GLY	A	108	-42.639	92.462	-30.375	1.00	32.12	T7
20	ATOM	7724	O	GLY	A	108	-42.687	92.544	-31.596	1.00	34.48	T7
	ATOM	7725	N	ILE	A	109	-41.522	92.663	-29.686	1.00	29.06	T7
	ATOM	7726	CA	ILE	A	109	-40.260	92.985	-30.343	1.00	34.42	T7
	ATOM	7727	CB	ILE	A	109	-39.616	94.234	-29.729	1.00	29.44	T7
	ATOM	7728	CG2	ILE	A	109	-38.306	94.521	-30.408	1.00	22.79	T7
25	ATOM	7729	CG1	ILE	A	109	-40.554	95.427	-29.865	1.00	25.28	T7
	ATOM	7730	CD1	ILE	A	109	-40.103	96.637	-29.092	1.00	29.17	T7
	ATOM	7731	C	ILE	A	109	-39.293	91.825	-30.180	1.00	29.40	T7
	ATOM	7732	O	ILE	A	109	-39.239	91.196	-29.128	1.00	25.62	T7
	ATOM	7733	N	ALA	A	110	-38.527	91.542	-31.222	1.00	32.13	T7
30	ATOM	7734	CA	ALA	A	110	-37.563	90.452	-31.177	1.00	26.69	T7
	ATOM	7735	CB	ALA	A	110	-38.202	89.173	-31.648	1.00	29.86	T7
	ATOM	7736	C	ALA	A	110	-36.381	90.786	-32.063	1.00	32.12	T7
	ATOM	7737	O	ALA	A	110	-36.506	91.573	-32.993	1.00	30.95	T7
	ATOM	7738	N	LYS	A	111	-35.227	90.201	-31.771	1.00	38.26	T7
35	ATOM	7739	CA	LYS	A	111	-34.059	90.457	-32.585	1.00	34.51	T7
	ATOM	7740	CB	LYS	A	111	-32.794	90.491	-31.744	1.00	34.95	T7
	ATOM	7741	CG	LYS	A	111	-31.570	90.810	-32.585	1.00	27.68	T7
	ATOM	7742	CD	LYS	A	111	-30.301	90.909	-31.749	1.00	26.50	T7
	ATOM	7743	CE	LYS	A	111	-29.081	91.269	-32.617	1.00	31.73	T7
40	ATOM	7744	NZ	LYS	A	111	-27.817	91.405	-31.798	1.00	29.30	T7
	ATOM	7745	C	LYS	A	111	-33.975	89.332	-33.587	1.00	33.72	T7
	ATOM	7746	O	LYS	A	111	-34.091	88.168	-33.214	1.00	35.08	T7
	ATOM	7747	N	LEU	A	112	-33.784	89.682	-34.858	1.00	33.89	T7
	ATOM	7748	CA	LEU	A	112	-33.710	88.701	-35.940	1.00	26.70	T7
45	ATOM	7749	CB	LEU	A	112	-34.926	88.841	-36.849	1.00	31.94	T7
	ATOM	7750	CG	LEU	A	112	-36.286	88.887	-36.168	1.00	34.92	T7
	ATOM	7751	CD1	LEU	A	112	-37.343	89.333	-37.150	1.00	28.18	T7
	ATOM	7752	CD2	LEU	A	112	-36.600	87.530	-35.599	1.00	28.61	T7
	ATOM	7753	C	LEU	A	112	-32.453	88.909	-36.775	1.00	24.28	T7
50	ATOM	7754	O	LEU	A	112	-31.855	89.982	-36.731	1.00	27.08	T7
	ATOM	7755	N	GLU	A	113	-32.068	87.886	-37.540	1.00	27.55	T7
	ATOM	7756	CA	GLU	A	113	-30.885	87.956	-38.395	1.00	33.64	T7
	ATOM	7757	CB	GLU	A	113	-29.797	87.018	-37.914	1.00	33.60	T7
	ATOM	7758	CG	GLU	A	113	-29.603	86.932	-36.442	1.00	34.90	T7
55	ATOM	7759	CD	GLU	A	113	-28.236	86.371	-36.112	1.00	26.96	T7
	ATOM	7760	OE1	GLU	A	113	-27.812	85.377	-36.772	1.00	34.94	T7
	ATOM	7761	OE2	GLU	A	113	-27.578	86.928	-35.193	1.00	30.86	T7
	ATOM	7762	C	GLU	A	113	-31.173	87.562	-39.825	1.00	28.96	T7
	ATOM	7763	O	GLU	A	113	-32.129	86.842	-40.095	1.00	35.95	T7
60	ATOM	7764	N	GLU	A	114	-30.317	88.019	-40.736	1.00	32.34	T7
	ATOM	7765	CA	GLU	A	114	-30.443	87.691	-42.153	1.00	32.17	T7
	ATOM	7766	CB	GLU	A	114	-29.137	87.940	-42.880	1.00	29.08	T7
	ATOM	7767	CG	GLU	A	114	-28.935	89.302	-43.437	1.00	31.76	T7
	ATOM	7768	CD	GLU	A	114	-27.864	89.277	-44.501	1.00	29.87	T7
65	ATOM	7769	OE1	GLU	A	114	-28.081	88.607	-45.545	1.00	35.87	T7
	ATOM	7770	OE2	GLU	A	114	-26.807	89.914	-44.286	1.00	33.64	T7

	ATOM	7771	C	GLU A 114	-30.730	86.217	-42.310	1.00	35.52	T7
	ATOM	7772	O	GLU A 114	-29.974	85.392	-41.819	1.00	33.91	T7
	ATOM	7773	N	GLY A 115	-31.800	85.877	-43.007	1.00	30.87	T7
	ATOM	7774	CA	GLY A 115	-32.092	84.475	-43.205	1.00	25.48	T7
5	ATOM	7775	C	GLY A 115	-33.143	83.915	-42.280	1.00	32.15	T7
	ATOM	7776	O	GLY A 115	-33.675	82.846	-42.540	1.00	27.59	T7
	ATOM	7777	N	ASP A 116	-33.432	84.600	-41.183	1.00	31.97	T7
	ATOM	7778	CA	ASP A 116	-34.457	84.097	-40.285	1.00	28.55	T7
	ATOM	7779	CB	ASP A 116	-34.525	84.920	-38.992	1.00	24.43	T7
10	ATOM	7780	CG	ASP A 116	-33.334	84.696	-38.082	1.00	28.28	T7
	ATOM	7781	OD1	ASP A 116	-32.721	83.608	-38.143	1.00	28.64	T7
	ATOM	7782	OD2	ASP A 116	-33.025	85.603	-37.285	1.00	35.08	T7
	ATOM	7783	C	ASP A 116	-35.786	84.209	-41.018	1.00	36.38	T7
	ATOM	7784	O	ASP A 116	-35.948	85.055	-41.902	1.00	31.96	T7
15	ATOM	7785	N	GLU A 117	-36.733	83.350	-40.663	1.00	29.34	T7
	ATOM	7786	CA	GLU A 117	-38.048	83.404	-41.283	1.00	31.02	T7
	ATOM	7787	CB	GLU A 117	-38.295	82.168	-42.136	1.00	32.04	T7
	ATOM	7788	CG	GLU A 117	-37.221	81.910	-43.156	1.00	31.80	T7
	ATOM	7789	CD	GLU A 117	-37.543	80.722	-44.037	1.00	29.76	T7
20	ATOM	7790	OE1	GLU A 117	-38.100	79.723	-43.516	1.00	29.76	T7
	ATOM	7791	OE2	GLU A 117	-37.226	80.785	-45.250	1.00	36.89	T7
	ATOM	7792	C	GLU A 117	-39.099	83.478	-40.187	1.00	31.28	T7
	ATOM	7793	O	GLU A 117	-38.962	82.845	-39.140	1.00	25.33	T7
	ATOM	7794	N	LEU A 118	-40.133	84.275	-40.420	1.00	33.44	T7
25	ATOM	7795	CA	LEU A 118	-41.212	84.416	-39.465	1.00	29.70	T7
	ATOM	7796	CB	LEU A 118	-41.523	85.885	-39.217	1.00	39.50	T7
	ATOM	7797	CG	LEU A 118	-40.448	86.718	-38.534	1.00	31.33	T7
	ATOM	7798	CD1	LEU A 118	-40.962	88.116	-38.297	1.00	38.34	T7
	ATOM	7799	CD2	LEU A 118	-40.062	86.072	-37.227	1.00	33.88	T7
30	ATOM	7800	C	LEU A 118	-42.439	83.754	-40.049	1.00	30.46	T7
	ATOM	7801	O	LEU A 118	-42.650	83.796	-41.256	1.00	28.07	T7
	ATOM	7802	N	GLN A 119	-43.250	83.141	-39.193	1.00	25.64	T7
	ATOM	7803	CA	GLN A 119	-44.475	82.493	-39.642	1.00	24.86	T7
	ATOM	7804	CB	GLN A 119	-44.178	81.064	-40.062	1.00	36.48	T7
35	ATOM	7805	CG	GLN A 119	-43.681	80.196	-38.939	1.00	31.83	T7
	ATOM	7806	CD	GLN A 119	-43.316	78.803	-39.409	1.00	26.20	T7
	ATOM	7807	OE1	GLN A 119	-43.204	77.875	-38.609	1.00	36.82	T7
	ATOM	7808	NE2	GLN A 119	-43.114	78.654	-40.711	1.00	36.83	T7
	ATOM	7809	C	GLN A 119	-45.555	82.517	-38.559	1.00	29.75	T7
40	ATOM	7810	O	GLN A 119	-45.254	82.610	-37.375	1.00	36.60	T7
	ATOM	7811	N	LEU A 120	-46.809	82.447	-38.990	1.00	30.49	T7
	ATOM	7812	CA	LEU A 120	-47.973	82.470	-38.108	1.00	30.76	T7
	ATOM	7813	CB	LEU A 120	-48.976	83.503	-38.633	1.00	30.89	T7
	ATOM	7814	CG	LEU A 120	-50.193	84.026	-37.871	1.00	32.90	T7
45	ATOM	7815	CD1	LEU A 120	-50.938	82.885	-37.234	1.00	34.53	T7
	ATOM	7816	CD2	LEU A 120	-49.740	85.016	-36.843	1.00	36.97	T7
	ATOM	7817	C	LEU A 120	-48.593	81.079	-38.156	1.00	31.87	T7
	ATOM	7818	O	LEU A 120	-48.997	80.613	-39.220	1.00	24.09	T7
	ATOM	7819	N	ALA A 121	-48.689	80.413	-37.017	1.00	37.91	T7
50	ATOM	7820	CA	ALA A 121	-49.244	79.070	-37.005	1.00	25.05	T7
	ATOM	7821	CB	ALA A 121	-48.132	78.077	-36.770	1.00	25.63	T7
	ATOM	7822	C	ALA A 121	-50.368	78.816	-36.008	1.00	35.78	T7
	ATOM	7823	O	ALA A 121	-50.341	79.300	-34.878	1.00	23.97	T7
	ATOM	7824	N	ILE A 122	-51.359	78.041	-36.434	1.00	36.47	T7
55	ATOM	7825	CA	ILE A 122	-52.480	77.689	-35.571	1.00	26.53	T7
	ATOM	7826	CB	ILE A 122	-53.821	77.895	-36.276	1.00	31.72	T7
	ATOM	7827	CG2	ILE A 122	-54.947	77.554	-35.326	1.00	31.10	T7
	ATOM	7828	CG1	ILE A 122	-53.943	79.347	-36.743	1.00	25.23	T7
	ATOM	7829	CD1	ILE A 122	-55.226	79.662	-37.455	1.00	29.47	T7
60	ATOM	7830	C	ILE A 122	-52.339	76.220	-35.188	1.00	30.04	T7
	ATOM	7831	O	ILE A 122	-52.360	75.340	-36.046	1.00	32.51	T7
	ATOM	7832	N	PRO A 123	-52.183	75.940	-33.885	1.00	29.11	T7
	ATOM	7833	CD	PRO A 123	-52.040	76.940	-32.818	1.00	25.11	T7
	ATOM	7834	CA	PRO A 123	-52.027	74.585	-33.340	1.00	28.30	T7
65	ATOM	7835	CB	PRO A 123	-51.587	74.833	-31.894	1.00	24.67	T7
	ATOM	7836	CG	PRO A 123	-51.082	76.248	-31.901	1.00	31.94	T7

	ATOM	7837	C	PRO	A	123	-53.304	73.742	-33.388	1.00	27.54	T7
	ATOM	7838	O	PRO	A	123	-53.745	73.232	-32.353	1.00	25.16	T7
	ATOM	7839	N	ARG	A	124	-53.900	73.606	-34.571	1.00	26.42	T7
	ATOM	7840	CA	ARG	A	124	-55.110	72.810	-34.739	1.00	35.02	T7
5	ATOM	7841	CB	ARG	A	124	-56.352	73.660	-34.582	1.00	30.43	T7
	ATOM	7842	CG	ARG	A	124	-56.793	73.785	-33.159	1.00	31.18	T7
	ATOM	7843	CD	ARG	A	124	-58.297	73.609	-33.077	1.00	37.22	T7
	ATOM	7844	NE	ARG	A	124	-58.709	72.220	-33.281	1.00	28.77	T7
	ATOM	7845	CZ	ARG	A	124	-59.974	71.843	-33.443	1.00	26.81	T7
10	ATOM	7846	NH1	ARG	A	124	-60.937	72.754	-33.427	1.00	39.27	T7
	ATOM	7847	NH2	ARG	A	124	-60.280	70.561	-33.614	1.00	30.26	T7
	ATOM	7848	C	ARG	A	124	-55.105	72.207	-36.111	1.00	32.66	T7
	ATOM	7849	O	ARG	A	124	-54.464	72.742	-37.013	1.00	30.99	T7
	ATOM	7850	N	GLU	A	125	-55.833	71.108	-36.280	1.00	28.17	T7
15	ATOM	7851	CA	GLU	A	125	-55.850	70.442	-37.570	1.00	31.21	T7
	ATOM	7852	CB	GLU	A	125	-56.355	69.016	-37.423	1.00	33.85	T7
	ATOM	7853	CG	GLU	A	125	-55.344	68.115	-36.726	1.00	31.72	T7
	ATOM	7854	CD	GLU	A	125	-55.264	66.740	-37.366	1.00	24.38	T7
	ATOM	7855	OE1	GLU	A	125	-56.328	66.066	-37.462	1.00	28.47	T7
20	ATOM	7856	OE2	GLU	A	125	-54.136	66.339	-37.768	1.00	32.90	T7
	ATOM	7857	C	GLU	A	125	-56.602	71.161	-38.670	1.00	28.74	T7
	ATOM	7858	O	GLU	A	125	-56.067	71.327	-39.769	1.00	36.65	T7
	ATOM	7859	N	ASN	A	126	-57.835	71.573	-38.414	1.00	31.27	T7
	ATOM	7860	CA	ASN	A	126	-58.556	72.307	-39.445	1.00	29.99	T7
25	ATOM	7861	CB	ASN	A	126	-59.549	71.415	-40.184	1.00	27.00	T7
	ATOM	7862	CG	ASN	A	126	-58.882	70.580	-41.265	1.00	34.68	T7
	ATOM	7863	OD1	ASN	A	126	-58.253	69.561	-40.975	1.00	33.30	T7
	ATOM	7864	ND2	ASN	A	126	-58.999	71.022	-42.519	1.00	29.25	T7
	ATOM	7865	C	ASN	A	126	-59.265	73.465	-38.803	1.00	39.91	T7
30	ATOM	7866	O	ASN	A	126	-60.495	73.550	-38.804	1.00	31.77	T7
	ATOM	7867	N	ALA	A	127	-58.457	74.359	-38.245	1.00	28.34	T7
	ATOM	7868	CA	ALA	A	127	-58.961	75.529	-37.559	1.00	28.92	T7
	ATOM	7869	CB	ALA	A	127	-57.811	76.473	-37.261	1.00	33.84	T7
	ATOM	7870	C	ALA	A	127	-60.033	76.247	-38.363	1.00	28.25	T7
35	ATOM	7871	O	ALA	A	127	-59.863	76.485	-39.554	1.00	29.14	T7
	ATOM	7872	N	GLN	A	128	-61.148	76.561	-37.707	1.00	34.83	T7
	ATOM	7873	CA	GLN	A	128	-62.225	77.303	-38.351	1.00	28.46	T7
	ATOM	7874	CB	GLN	A	128	-63.550	77.008	-37.658	1.00	25.89	T7
	ATOM	7875	CG	GLN	A	128	-63.992	75.567	-37.834	1.00	31.95	T7
40	ATOM	7876	CD	GLN	A	128	-63.870	75.112	-39.274	1.00	32.41	T7
	ATOM	7877	OE1	GLN	A	128	-64.473	75.692	-40.175	1.00	29.65	T7
	ATOM	7878	NE2	GLN	A	128	-63.075	74.074	-39.497	1.00	31.04	T7
	ATOM	7879	C	GLN	A	128	-61.845	78.780	-38.222	1.00	27.73	T7
	ATOM	7880	O	GLN	A	128	-62.037	79.411	-37.181	1.00	26.22	T7
45	ATOM	7881	N	ILE	A	129	-61.306	79.319	-39.304	1.00	34.47	T7
	ATOM	7882	CA	ILE	A	129	-60.800	80.678	-39.342	1.00	35.46	T7
	ATOM	7883	CB	ILE	A	129	-59.260	80.587	-39.606	1.00	38.11	T7
	ATOM	7884	CG2	ILE	A	129	-58.826	81.456	-40.748	1.00	32.20	T7
	ATOM	7885	CG1	ILE	A	129	-58.513	80.886	-38.326	1.00	27.30	T7
50	ATOM	7886	CD1	ILE	A	129	-58.851	79.924	-37.227	1.00	43.12	T7
	ATOM	7887	C	ILE	A	129	-61.477	81.559	-40.386	1.00	28.98	T7
	ATOM	7888	O	ILE	A	129	-62.082	81.058	-41.327	1.00	27.51	T7
	ATOM	7889	N	SER	A	130	-61.395	82.872	-40.211	1.00	24.44	T7
	ATOM	7890	CA	SER	A	130	-61.956	83.794	-41.193	1.00	31.80	T7
55	ATOM	7891	CB	SER	A	130	-62.640	84.971	-40.515	1.00	37.83	T7
	ATOM	7892	OG	SER	A	130	-62.934	85.984	-41.468	1.00	33.26	T7
	ATOM	7893	C	SER	A	130	-60.795	84.318	-42.020	1.00	30.65	T7
	ATOM	7894	O	SER	A	130	-59.832	84.824	-41.462	1.00	34.74	T7
	ATOM	7895	N	LEU	A	131	-60.875	84.208	-43.341	1.00	27.75	T7
60	ATOM	7896	CA	LEU	A	131	-59.784	84.676	-44.188	1.00	30.93	T7
	ATOM	7897	CB	LEU	A	131	-59.510	83.685	-45.317	1.00	29.92	T7
	ATOM	7898	CG	LEU	A	131	-58.700	82.447	-44.936	1.00	32.26	T7
	ATOM	7899	CD1	LEU	A	131	-59.399	81.671	-43.866	1.00	31.61	T7
	ATOM	7900	CD2	LEU	A	131	-58.520	81.584	-46.146	1.00	28.04	T7
65	ATOM	7901	C	LEU	A	131	-59.990	86.055	-44.777	1.00	34.12	T7
	ATOM	7902	O	LEU	A	131	-59.546	86.331	-45.890	1.00	31.15	T7

	ATOM	7903	N	ASP A 132	-60.653	86.929	-44.030	1.00	27.49	T7
	ATOM	7904	CA	ASP A 132	-60.886	88.293	-44.497	1.00	33.70	T7
	ATOM	7905	CB	ASP A 132	-62.177	88.857	-43.886	1.00	27.75	T7
	ATOM	7906	CG	ASP A 132	-63.424	88.424	-44.647	1.00	28.41	T7
5	ATOM	7907	OD1	ASP A 132	-64.543	88.629	-44.115	1.00	30.03	T7
	ATOM	7908	OD2	ASP A 132	-63.282	87.894	-45.776	1.00	33.89	T7
	ATOM	7909	C	ASP A 132	-59.713	89.221	-44.177	1.00	29.16	T7
	ATOM	7910	O	ASP A 132	-59.103	89.136	-43.111	1.00	35.25	T7
	ATOM	7911	N	GLY A 133	-59.413	90.105	-45.121	1.00	31.35	T7
10	ATOM	7912	CA	GLY A 133	-58.326	91.043	-44.952	1.00	33.68	T7
	ATOM	7913	C	GLY A 133	-58.366	91.870	-43.682	1.00	28.83	T7
	ATOM	7914	O	GLY A 133	-57.313	92.217	-43.172	1.00	32.07	T7
	ATOM	7915	N	ASP A 134	-59.553	92.195	-43.170	1.00	37.12	T7
	ATOM	7916	CA	ASP A 134	-59.652	92.997	-41.951	1.00	34.19	T7
15	ATOM	7917	CB	ASP A 134	-61.078	93.423	-41.621	1.00	23.31	T7
	ATOM	7918	CG	ASP A 134	-61.956	93.491	-42.799	1.00	28.05	T7
	ATOM	7919	OD1	ASP A 134	-61.653	94.317	-43.667	1.00	20.40	T7
	ATOM	7920	OD2	ASP A 134	-62.948	92.733	-42.847	1.00	33.80	T7
	ATOM	7921	C	ASP A 134	-59.238	92.214	-40.741	1.00	28.20	T7
20	ATOM	7922	O	ASP A 134	-58.354	92.603	-40.001	1.00	31.18	T7
	ATOM	7923	N	VAL A 135	-59.936	91.112	-40.531	1.00	37.48	T7
	ATOM	7924	CA	VAL A 135	-59.729	90.282	-39.375	1.00	34.28	T7
	ATOM	7925	CB	VAL A 135	-60.900	89.343	-39.246	1.00	30.68	T7
	ATOM	7926	CG1	VAL A 135	-62.163	90.160	-39.091	1.00	26.52	T7
25	ATOM	7927	CG2	VAL A 135	-60.993	88.467	-40.473	1.00	24.91	T7
	ATOM	7928	C	VAL A 135	-58.433	89.510	-39.178	1.00	26.61	T7
	ATOM	7929	O	VAL A 135	-57.977	89.407	-38.046	1.00	29.56	T7
	ATOM	7930	N	THR A 136	-57.827	88.959	-40.228	1.00	28.31	T7
	ATOM	7931	CA	THR A 136	-56.573	88.233	-40.005	1.00	28.41	T7
30	ATOM	7932	CB	THR A 136	-56.765	86.700	-40.158	1.00	24.59	T7
	ATOM	7933	OG1	THR A 136	-56.800	86.357	-41.534	1.00	28.17	T7
	ATOM	7934	CG2	THR A 136	-58.074	86.259	-39.528	1.00	30.42	T7
	ATOM	7935	C	THR A 136	-55.412	88.719	-40.882	1.00	29.36	T7
	ATOM	7936	O	THR A 136	-55.481	88.696	-42.109	1.00	34.25	T7
35	ATOM	7937	N	PHE A 137	-54.345	89.171	-40.221	1.00	34.17	T7
	ATOM	7938	CA	PHE A 137	-53.163	89.701	-40.899	1.00	27.84	T7
	ATOM	7939	CB	PHE A 137	-53.326	91.210	-41.082	1.00	29.10	T7
	ATOM	7940	CG	PHE A 137	-53.858	91.917	-39.868	1.00	26.72	T7
	ATOM	7941	CD1	PHE A 137	-53.021	92.260	-38.825	1.00	30.29	T7
40	ATOM	7942	CD2	PHE A 137	-55.193	92.246	-39.774	1.00	27.50	T7
	ATOM	7943	CE1	PHE A 137	-53.508	92.918	-37.711	1.00	24.72	T7
	ATOM	7944	CE2	PHE A 137	-55.689	92.902	-38.664	1.00	32.47	T7
	ATOM	7945	CZ	PHE A 137	-54.847	93.239	-37.635	1.00	33.56	T7
	ATOM	7946	C	PHE A 137	-51.864	89.385	-40.160	1.00	32.84	T7
45	ATOM	7947	O	PHE A 137	-51.887	89.012	-38.989	1.00	33.22	T7
	ATOM	7948	N	PHE A 138	-50.732	89.545	-40.843	1.00	27.31	T7
	ATOM	7949	CA	PHE A 138	-49.434	89.228	-40.253	1.00	24.78	T7
	ATOM	7950	CB	PHE A 138	-48.915	87.949	-40.913	1.00	28.35	T7
	ATOM	7951	CG	PHE A 138	-47.763	87.312	-40.204	1.00	33.35	T7
50	ATOM	7952	CD1	PHE A 138	-47.531	87.546	-38.857	1.00	30.81	T7
	ATOM	7953	CD2	PHE A 138	-46.902	86.469	-40.895	1.00	27.42	T7
	ATOM	7954	CE1	PHE A 138	-46.451	86.952	-38.211	1.00	35.81	T7
	ATOM	7955	CE2	PHE A 138	-45.822	85.872	-40.259	1.00	41.84	T7
	ATOM	7956	CZ	PHE A 138	-45.596	86.115	-38.917	1.00	28.31	T7
55	ATOM	7957	C	PHE A 138	-48.435	90.392	-40.362	1.00	22.75	T7
	ATOM	7958	O	PHE A 138	-48.234	90.962	-41.426	1.00	39.78	T7
	ATOM	7959	N	GLY A 139	-47.809	90.699	-39.222	1.00	35.40	T7
	ATOM	7960	CA	GLY A 139	-46.892	91.825	-39.053	1.00	31.03	T7
	ATOM	7961	C	GLY A 139	-45.492	91.972	-39.595	1.00	36.11	T7
60	ATOM	7962	O	GLY A 139	-45.268	91.720	-40.762	1.00	32.35	T7
	ATOM	7963	N	ALA A 140	-44.579	92.445	-38.744	1.00	30.48	T7
	ATOM	7964	CA	ALA A 140	-43.153	92.683	-39.061	1.00	29.25	T7
	ATOM	7965	CB	ALA A 140	-42.636	91.627	-40.015	1.00	30.89	T7
	ATOM	7966	C	ALA A 140	-42.759	94.074	-39.587	1.00	24.68	T7
65	ATOM	7967	O	ALA A 140	-42.989	94.408	-40.737	1.00	28.92	T7
	ATOM	7968	N	LEU A 141	-42.138	94.864	-38.718	1.00	31.99	T7

	ATOM	7969	CA	LEU	A	141	-41.669	96.213	-39.037	1.00	31.68	T7
	ATOM	7970	CB	LEU	A	141	-42.654	97.258	-38.498	1.00	29.07	T7
	ATOM	7971	CG	LEU	A	141	-42.277	98.745	-38.554	1.00	31.81	T7
	ATOM	7972	CD1	LEU	A	141	-43.498	99.596	-38.390	1.00	27.65	T7
5	ATOM	7973	CD2	LEU	A	141	-41.292	99.070	-37.466	1.00	31.16	T7
	ATOM	7974	C	LEU	A	141	-40.297	96.403	-38.382	1.00	27.31	T7
	ATOM	7975	O	LEU	A	141	-40.113	96.066	-37.217	1.00	32.75	T7
	ATOM	7976	N	LYS	A	142	-39.337	96.954	-39.118	1.00	31.24	T7
	ATOM	7977	CA	LYS	A	142	-38.001	97.141	-38.565	1.00	35.17	T7
10	ATOM	7978	CB	LYS	A	142	-36.959	97.030	-39.672	1.00	28.75	T7
	ATOM	7979	CG	LYS	A	142	-35.550	97.218	-39.158	1.00	34.06	T7
	ATOM	7980	CD	LYS	A	142	-34.500	96.835	-40.176	1.00	34.23	T7
	ATOM	7981	CE	LYS	A	142	-33.114	97.017	-39.581	1.00	30.03	T7
	ATOM	7982	NZ	LYS	A	142	-32.035	96.674	-40.549	1.00	27.87	T7
15	ATOM	7983	C	LYS	A	142	-37.770	98.440	-37.785	1.00	27.80	T7
	ATOM	7984	O	LYS	A	142	-38.050	99.535	-38.271	1.00	29.50	T7
	ATOM	7985	N	LEU	A	143	-37.241	98.310	-36.573	1.00	35.29	T7
	ATOM	7986	CA	LEU	A	143	-36.973	99.470	-35.730	1.00	20.87	T7
	ATOM	7987	CB	LEU	A	143	-36.907	99.048	-34.260	1.00	25.71	T7
20	ATOM	7988	CG	LEU	A	143	-38.119	98.359	-33.631	1.00	35.23	T7
	ATOM	7989	CD1	LEU	A	143	-37.783	97.946	-32.209	1.00	26.94	T7
	ATOM	7990	CD2	LEU	A	143	-39.308	99.294	-33.641	1.00	25.05	T7
	ATOM	7991	C	LEU	A	143	-35.646	100.107	-36.119	1.00	37.35	T7
	ATOM	7992	O	LEU	A	143	-34.774	99.442	-36.673	1.00	28.91	T7
25	ATOM	7993	N	LEU	A	144	-35.488	101.391	-35.822	1.00	26.39	T7
	ATOM	7994	CA	LEU	A	144	-34.248	102.086	-36.135	1.00	28.03	T7
	ATOM	7995	CB	LEU	A	144	-34.504	103.572	-36.365	1.00	30.82	T7
	ATOM	7996	CG	LEU	A	144	-35.360	103.920	-37.584	1.00	26.00	T7
	ATOM	7997	CD1	LEU	A	144	-35.675	105.406	-37.557	1.00	29.34	T7
30	ATOM	7998	CD2	LEU	A	144	-34.635	103.538	-38.876	1.00	29.26	T7
	ATOM	7999	C	LEU	A	144	-33.260	101.934	-34.994	1.00	37.66	T7
	ATOM	8000	O	LEU	A	144	-33.683	101.573	-33.877	1.00	30.90	T7
	ATOM	8001	OXT	LEU	A	144	-32.068	102.203	-35.237	1.00	31.68	T7
	ATOM	8002	CB	VAL	A	1	-28.829	107.420	-34.319	1.00	33.90	T8
35	ATOM	8003	CG1	VAL	A	1	-28.292	106.434	-35.375	1.00	29.35	T8
	ATOM	8004	CG2	VAL	A	1	-28.052	108.734	-34.383	1.00	30.26	T8
	ATOM	8005	C	VAL	A	1	-31.135	106.366	-34.357	1.00	27.19	T8
	ATOM	8006	O	VAL	A	1	-31.776	105.877	-35.305	1.00	27.52	T8
	ATOM	8007	N	VAL	A	1	-30.865	108.772	-33.654	1.00	25.41	T8
40	ATOM	8008	CA	VAL	A	1	-30.360	107.689	-34.551	1.00	27.22	T8
	ATOM	8009	N	THR	A	2	-31.089	105.789	-33.150	1.00	27.31	T8
	ATOM	8010	CA	THR	A	2	-31.789	104.528	-32.908	1.00	24.16	T8
	ATOM	8011	CB	THR	A	2	-30.849	103.452	-32.353	1.00	31.25	T8
	ATOM	8012	OG1	THR	A	2	-30.510	103.781	-31.002	1.00	29.13	T8
45	ATOM	8013	CG2	THR	A	2	-29.579	103.352	-33.207	1.00	28.29	T8
	ATOM	8014	C	THR	A	2	-32.945	104.671	-31.931	1.00	25.06	T8
	ATOM	8015	O	THR	A	2	-33.121	105.721	-31.319	1.00	31.83	T8
	ATOM	8016	N	GLN	A	3	-33.726	103.600	-31.785	1.00	30.47	T8
	ATOM	8017	CA	GLN	A	3	-34.877	103.596	-30.880	1.00	30.46	T8
50	ATOM	8018	CB	GLN	A	3	-36.138	103.166	-31.629	1.00	32.37	T8
	ATOM	8019	CG	GLN	A	3	-36.330	103.838	-32.962	1.00	36.69	T8
	ATOM	8020	CD	GLN	A	3	-37.664	103.495	-33.577	1.00	25.27	T8
	ATOM	8021	OE1	GLN	A	3	-38.707	103.790	-32.996	1.00	24.61	T8
	ATOM	8022	NE2	GLN	A	3	-37.645	102.871	-34.755	1.00	29.03	T8
55	ATOM	8023	C	GLN	A	3	-34.669	102.648	-29.703	1.00	32.84	T8
	ATOM	8024	O	GLN	A	3	-34.776	101.432	-29.854	1.00	35.83	T8
	ATOM	8025	N	ASP	A	4	-34.380	103.196	-28.532	1.00	35.22	T8
	ATOM	8026	CA	ASP	A	4	-34.185	102.350	-27.372	1.00	30.04	T8
	ATOM	8027	CB	ASP	A	4	-33.799	103.190	-26.161	1.00	37.16	T8
60	ATOM	8028	CG	ASP	A	4	-32.474	103.895	-26.340	1.00	27.06	T8
	ATOM	8029	OD1	ASP	A	4	-31.714	103.487	-27.243	1.00	37.09	T8
	ATOM	8030	OD2	ASP	A	4	-32.186	104.842	-25.573	1.00	35.07	T8
	ATOM	8031	C	ASP	A	4	-35.460	101.583	-27.066	1.00	28.85	T8
	ATOM	8032	O	ASP	A	4	-36.558	102.072	-27.307	1.00	29.92	T8
65	ATOM	8033	N	CYS	A	5	-35.305	100.372	-26.547	1.00	35.09	T8
	ATOM	8034	CA	CYS	A	5	-36.436	99.522	-26.183	1.00	33.41	T8

	ATOM	8035	CB	CYS	A	5	-37.010	98.796	-27.418	1.00	29.01	T8
	ATOM	8036	SG	CYS	A	5	-35.847	98.391	-28.734	1.00	36.92	T8
	ATOM	8037	C	CYS	A	5	-36.006	98.507	-25.129	1.00	31.42	T8
	ATOM	8038	O	CYS	A	5	-34.861	98.074	-25.099	1.00	37.47	T8
5	ATOM	8039	N	LEU	A	6	-36.921	98.154	-24.240	1.00	33.63	T8
	ATOM	8040	CA	LEU	A	6	-36.632	97.172	-23.203	1.00	28.27	T8
	ATOM	8041	CB	LEU	A	6	-36.361	97.863	-21.869	1.00	33.43	T8
	ATOM	8042	CG	LEU	A	6	-36.101	96.959	-20.668	1.00	28.44	T8
	ATOM	8043	CD1	LEU	A	6	-35.191	97.657	-19.698	1.00	28.76	T8
10	ATOM	8044	CD2	LEU	A	6	-37.405	96.603	-20.006	1.00	29.82	T8
	ATOM	8045	C	LEU	A	6	-37.850	96.270	-23.097	1.00	28.66	T8
	ATOM	8046	O	LEU	A	6	-38.977	96.751	-23.080	1.00	29.28	T8
	ATOM	8047	N	GLN	A	7	-37.630	94.962	-23.042	1.00	33.41	T8
	ATOM	8048	CA	GLN	A	7	-38.742	94.025	-22.961	1.00	34.77	T8
15	ATOM	8049	CB	GLN	A	7	-38.890	93.291	-24.286	1.00	26.94	T8
	ATOM	8050	CG	GLN	A	7	-40.152	92.486	-24.419	1.00	28.72	T8
	ATOM	8051	CD	GLN	A	7	-40.343	91.979	-25.823	1.00	41.43	T8
	ATOM	8052	OE1	GLN	A	7	-39.620	91.110	-26.277	1.00	32.77	T8
	ATOM	8053	NE2	GLN	A	7	-41.308	92.537	-26.526	1.00	33.06	T8
20	ATOM	8054	C	GLN	A	7	-38.571	93.017	-21.835	1.00	30.77	T8
	ATOM	8055	O	GLN	A	7	-37.471	92.531	-21.588	1.00	32.33	T8
	ATOM	8056	N	LEU	A	8	-39.672	92.712	-21.157	1.00	34.33	T8
	ATOM	8057	CA	LEU	A	8	-39.662	91.769	-20.060	1.00	30.40	T8
	ATOM	8058	CB	LEU	A	8	-40.189	92.443	-18.800	1.00	34.17	T8
25	ATOM	8059	CG	LEU	A	8	-39.252	93.235	-17.884	1.00	33.41	T8
	ATOM	8060	CD1	LEU	A	8	-37.947	93.527	-18.573	1.00	29.15	T8
	ATOM	8061	CD2	LEU	A	8	-39.950	94.511	-17.449	1.00	26.10	T8
	ATOM	8062	C	LEU	A	8	-40.493	90.534	-20.371	1.00	25.08	T8
	ATOM	8063	O	LEU	A	8	-41.432	90.579	-21.154	1.00	35.42	T8
30	ATOM	8064	N	ILE	A	9	-40.131	89.429	-19.734	1.00	34.23	T8
	ATOM	8065	CA	ILE	A	9	-40.799	88.144	-19.905	1.00	28.84	T8
	ATOM	8066	CB	ILE	A	9	-39.855	87.147	-20.583	1.00	26.08	T8
	ATOM	8067	CG2	ILE	A	9	-40.320	85.734	-20.347	1.00	27.48	T8
	ATOM	8068	CG1	ILE	A	9	-39.780	87.414	-22.069	1.00	35.41	T8
35	ATOM	8069	CD1	ILE	A	9	-38.838	86.458	-22.746	1.00	34.51	T8
	ATOM	8070	C	ILE	A	9	-41.159	87.565	-18.541	1.00	33.49	T8
	ATOM	8071	O	ILE	A	9	-40.405	87.719	-17.587	1.00	32.68	T8
	ATOM	8072	N	ALA	A	10	-42.291	86.880	-18.445	1.00	27.48	T8
	ATOM	8073	CA	ALA	A	10	-42.681	86.287	-17.172	1.00	31.73	T8
40	ATOM	8074	CB	ALA	A	10	-44.031	85.631	-17.288	1.00	27.31	T8
	ATOM	8075	C	ALA	A	10	-41.652	85.258	-16.718	1.00	37.57	T8
	ATOM	8076	O	ALA	A	10	-41.232	84.393	-17.492	1.00	26.23	T8
	ATOM	8077	N	ASP	A	11	-41.241	85.361	-15.458	1.00	35.03	T8
	ATOM	8078	CA	ASP	A	11	-40.273	84.429	-14.896	1.00	31.16	T8
45	ATOM	8079	CB	ASP	A	11	-39.381	85.122	-13.869	1.00	33.35	T8
	ATOM	8080	CG	ASP	A	11	-38.459	84.151	-13.165	1.00	33.90	T8
	ATOM	8081	OD1	ASP	A	11	-37.959	83.225	-13.847	1.00	31.70	T8
	ATOM	8082	OD2	ASP	A	11	-38.230	84.313	-11.945	1.00	38.20	T8
	ATOM	8083	C	ASP	A	11	-41.004	83.272	-14.240	1.00	25.93	T8
50	ATOM	8084	O	ASP	A	11	-41.341	83.318	-13.057	1.00	33.27	T8
	ATOM	8085	N	SER	A	12	-41.234	82.236	-15.038	1.00	32.29	T8
	ATOM	8086	CA	SER	A	12	-41.941	81.034	-14.617	1.00	20.97	T8
	ATOM	8087	CB	SER	A	12	-42.160	80.101	-15.820	1.00	26.99	T8
	ATOM	8088	OG	SER	A	12	-40.931	79.726	-16.434	1.00	27.84	T8
55	ATOM	8089	C	SER	A	12	-41.237	80.261	-13.519	1.00	28.72	T8
	ATOM	8090	O	SER	A	12	-41.485	79.073	-13.347	1.00	30.09	T8
	ATOM	8091	N	GLU	A	13	-40.362	80.921	-12.771	1.00	28.81	T8
	ATOM	8092	CA	GLU	A	13	-39.662	80.222	-11.708	1.00	30.44	T8
	ATOM	8093	CB	GLU	A	13	-38.281	79.818	-12.177	1.00	29.75	T8
60	ATOM	8094	CG	GLU	A	13	-38.299	78.462	-12.821	1.00	33.97	T8
	ATOM	8095	CD	GLU	A	13	-36.919	78.014	-13.211	1.00	26.81	T8
	ATOM	8096	OE1	GLU	A	13	-35.984	78.219	-12.382	1.00	27.39	T8
	ATOM	8097	OE2	GLU	A	13	-36.765	77.454	-14.335	1.00	30.90	T8
	ATOM	8098	C	GLU	A	13	-39.569	80.931	-10.379	1.00	30.12	T8
65	ATOM	8099	O	GLU	A	13	-38.659	80.680	-9.593	1.00	37.56	T8
	ATOM	8100	N	THR	A	14	-40.511	81.831	-10.140	1.00	28.83	T8

	ATOM	8101	CA	THR	A	14	-40.592	82.545	-8.880	1.00	30.02	T8
	ATOM	8102	CB	THR	A	14	-39.817	83.877	-8.884	1.00	32.38	T8
	ATOM	8103	OG1	THR	A	14	-40.301	84.706	-9.938	1.00	27.14	T8
	ATOM	8104	CG2	THR	A	14	-38.328	83.636	-9.068	1.00	24.95	T8
5	ATOM	8105	C	THR	A	14	-42.072	82.805	-8.665	1.00	26.16	T8
	ATOM	8106	O	THR	A	14	-42.847	82.922	-9.618	1.00	33.95	T8
	ATOM	8107	N	PRO	A	15	-42.490	82.871	-7.404	1.00	26.94	T8
	ATOM	8108	CD	PRO	A	15	-41.654	82.758	-6.201	1.00	38.80	T8
	ATOM	8109	CA	PRO	A	15	-43.888	83.109	-7.054	1.00	33.38	T8
10	ATOM	8110	CB	PRO	A	15	-43.871	83.066	-5.528	1.00	28.14	T8
	ATOM	8111	CG	PRO	A	15	-42.638	82.251	-5.205	1.00	35.09	T8
	ATOM	8112	C	PRO	A	15	-44.362	84.455	-7.567	1.00	31.87	T8
	ATOM	8113	O	PRO	A	15	-43.591	85.406	-7.606	1.00	36.62	T8
	ATOM	8114	N	THR	A	16	-45.628	84.536	-7.952	1.00	34.66	T8
15	ATOM	8115	CA	THR	A	16	-46.177	85.797	-8.424	1.00	32.20	T8
	ATOM	8116	CB	THR	A	16	-47.540	85.581	-9.058	1.00	30.58	T8
	ATOM	8117	OG1	THR	A	16	-48.491	85.232	-8.046	1.00	27.93	T8
	ATOM	8118	CG2	THR	A	16	-47.462	84.454	-10.046	1.00	27.74	T8
	ATOM	8119	C	THR	A	16	-46.328	86.730	-7.216	1.00	33.21	T8
20	ATOM	8120	O	THR	A	16	-47.092	86.446	-6.294	1.00	30.58	T8
	ATOM	8121	N	ILE	A	17	-45.599	87.840	-7.221	1.00	34.31	T8
	ATOM	8122	CA	ILE	A	17	-45.637	88.794	-6.119	1.00	29.22	T8
	ATOM	8123	CB	ILE	A	17	-44.817	90.034	-6.466	1.00	26.76	T8
	ATOM	8124	CG2	ILE	A	17	-44.897	91.041	-5.340	1.00	28.94	T8
25	ATOM	8125	CG1	ILE	A	17	-43.368	89.626	-6.739	1.00	29.89	T8
	ATOM	8126	CD1	ILE	A	17	-42.475	90.762	-7.155	1.00	34.20	T8
	ATOM	8127	C	ILE	A	17	-47.031	89.248	-5.700	1.00	28.81	T8
	ATOM	8128	O	ILE	A	17	-47.851	89.632	-6.535	1.00	28.57	T8
	ATOM	8129	N	GLN	A	18	-47.291	89.207	-4.394	1.00	30.62	T8
30	ATOM	8130	CA	GLN	A	18	-48.578	89.630	-3.843	1.00	30.90	T8
	ATOM	8131	CB	GLN	A	18	-49.194	88.519	-3.006	1.00	33.66	T8
	ATOM	8132	CG	GLN	A	18	-50.575	88.157	-3.462	1.00	34.77	T8
	ATOM	8133	CD	GLN	A	18	-50.542	87.383	-4.744	1.00	33.47	T8
	ATOM	8134	OE1	GLN	A	18	-50.204	86.206	-4.758	1.00	25.25	T8
35	ATOM	8135	NE2	GLN	A	18	-50.874	88.041	-5.840	1.00	27.96	T8
	ATOM	8136	C	GLN	A	18	-48.417	90.882	-2.983	1.00	36.25	T8
	ATOM	8137	O	GLN	A	18	-47.507	90.963	-2.160	1.00	30.63	T8
	ATOM	8138	N	LYS	A	19	-49.303	91.855	-3.168	1.00	29.65	T8
	ATOM	8139	CA	LYS	A	19	-49.218	93.094	-2.408	1.00	31.48	T8
40	ATOM	8140	CB	LYS	A	19	-47.951	93.838	-2.789	1.00	33.00	T8
	ATOM	8141	CG	LYS	A	19	-47.890	95.242	-2.250	1.00	27.79	T8
	ATOM	8142	CD	LYS	A	19	-46.510	95.835	-2.449	1.00	24.74	T8
	ATOM	8143	CE	LYS	A	19	-46.425	97.268	-1.916	1.00	42.73	T8
	ATOM	8144	NZ	LYS	A	19	-45.051	97.876	-2.076	1.00	27.80	T8
45	ATOM	8145	C	LYS	A	19	-50.413	94.008	-2.611	1.00	29.64	T8
	ATOM	8146	O	LYS	A	19	-50.788	94.298	-3.745	1.00	31.26	T8
	ATOM	8147	N	GLY	A	20	-50.994	94.472	-1.506	1.00	31.51	T8
	ATOM	8148	CA	GLY	A	20	-52.158	95.340	-1.577	1.00	29.28	T8
	ATOM	8149	C	GLY	A	20	-53.262	94.689	-2.389	1.00	31.32	T8
50	ATOM	8150	O	GLY	A	20	-53.914	95.352	-3.194	1.00	29.15	T8
	ATOM	8151	N	SER	A	21	-53.457	93.386	-2.171	1.00	29.53	T8
	ATOM	8152	CA	SER	A	21	-54.459	92.593	-2.885	1.00	23.72	T8
	ATOM	8153	CB	SER	A	21	-55.869	92.921	-2.364	1.00	30.07	T8
	ATOM	8154	OG	SER	A	21	-56.196	94.286	-2.539	1.00	31.43	T8
55	ATOM	8155	C	SER	A	21	-54.375	92.772	-4.413	1.00	30.04	T8
	ATOM	8156	O	SER	A	21	-55.384	92.846	-5.116	1.00	32.13	T8
	ATOM	8157	N	TYR	A	22	-53.142	92.839	-4.902	1.00	31.55	T8
	ATOM	8158	CA	TYR	A	22	-52.834	92.982	-6.316	1.00	36.17	T8
	ATOM	8159	CB	TYR	A	22	-52.197	94.338	-6.587	1.00	36.58	T8
60	ATOM	8160	CG	TYR	A	22	-53.158	95.400	-7.043	1.00	29.68	T8
	ATOM	8161	CD1	TYR	A	22	-54.508	95.332	-6.717	1.00	29.32	T8
	ATOM	8162	CE1	TYR	A	22	-55.400	96.339	-7.110	1.00	32.23	T8
	ATOM	8163	CD2	TYR	A	22	-52.716	96.497	-7.772	1.00	29.75	T8
	ATOM	8164	CE2	TYR	A	22	-53.597	97.509	-8.164	1.00	33.01	T8
65	ATOM	8165	CZ	TYR	A	22	-54.937	97.422	-7.830	1.00	35.83	T8
	ATOM	8166	OH	TYR	A	22	-55.821	98.408	-8.206	1.00	28.36	T8

	ATOM	8167	C	TYR	A	22	-51.825	91.901	-6.643	1.00	31.81	T8
	ATOM	8168	O	TYR	A	22	-51.005	91.544	-5.795	1.00	29.71	T8
	ATOM	8169	N	THR	A	23	-51.884	91.370	-7.861	1.00	29.41	T8
	ATOM	8170	CA	THR	A	23	-50.932	90.345	-8.252	1.00	32.48	T8
5	ATOM	8171	CB	THR	A	23	-51.610	89.131	-8.910	1.00	31.75	T8
	ATOM	8172	OG1	THR	A	23	-52.815	88.802	-8.211	1.00	33.22	T8
	ATOM	8173	CG2	THR	A	23	-50.688	87.936	-8.837	1.00	30.65	T8
	ATOM	8174	C	THR	A	23	-49.954	90.958	-9.242	1.00	28.58	T8
	ATOM	8175	O	THR	A	23	-50.358	91.607	-10.203	1.00	31.89	T8
10	ATOM	8176	N	PHE	A	24	-48.664	90.763	-8.990	1.00	28.71	T8
	ATOM	8177	CA	PHE	A	24	-47.628	91.295	-9.860	1.00	31.38	T8
	ATOM	8178	CB	PHE	A	24	-46.723	92.253	-9.084	1.00	22.89	T8
	ATOM	8179	CG	PHE	A	24	-47.414	93.496	-8.624	1.00	24.77	T8
	ATOM	8180	CD1	PHE	A	24	-48.195	93.491	-7.479	1.00	30.68	T8
15	ATOM	8181	CD2	PHE	A	24	-47.318	94.666	-9.360	1.00	24.58	T8
	ATOM	8182	CE1	PHE	A	24	-48.872	94.635	-7.073	1.00	33.10	T8
	ATOM	8183	CE2	PHE	A	24	-47.989	95.810	-8.966	1.00	32.14	T8
	ATOM	8184	CZ	PHE	A	24	-48.769	95.793	-7.819	1.00	27.85	T8
	ATOM	8185	C	PHE	A	24	-46.787	90.186	-10.480	1.00	28.63	T8
20	ATOM	8186	O	PHE	A	24	-46.215	89.355	-9.778	1.00	32.15	T8
	ATOM	8187	N	VAL	A	25	-46.723	90.182	-11.805	1.00	34.00	T8
	ATOM	8188	CA	VAL	A	25	-45.960	89.187	-12.539	1.00	25.09	T8
	ATOM	8189	CB	VAL	A	25	-46.176	89.338	-14.056	1.00	26.46	T8
	ATOM	8190	CG1	VAL	A	25	-45.311	88.354	-14.809	1.00	29.29	T8
25	ATOM	8191	CG2	VAL	A	25	-47.634	89.129	-14.396	1.00	22.23	T8
	ATOM	8192	C	VAL	A	25	-44.475	89.330	-12.259	1.00	26.98	T8
	ATOM	8193	O	VAL	A	25	-43.939	90.429	-12.268	1.00	34.49	T8
	ATOM	8194	N	PRO	A	26	-43.792	88.213	-11.988	1.00	26.99	T8
	ATOM	8195	CD	PRO	A	26	-44.369	86.887	-11.735	1.00	26.73	T8
30	ATOM	8196	CA	PRO	A	26	-42.353	88.217	-11.711	1.00	36.31	T8
	ATOM	8197	CB	PRO	A	26	-42.109	86.837	-11.119	1.00	34.06	T8
	ATOM	8198	CG	PRO	A	26	-43.470	86.380	-10.672	1.00	29.03	T8
	ATOM	8199	C	PRO	A	26	-41.647	88.366	-13.054	1.00	29.86	T8
	ATOM	8200	O	PRO	A	26	-41.684	87.451	-13.870	1.00	28.88	T8
35	ATOM	8201	N	TRP	A	27	-41.001	89.498	-13.298	1.00	30.07	T8
	ATOM	8202	CA	TRP	A	27	-40.337	89.691	-14.583	1.00	29.42	T8
	ATOM	8203	CB	TRP	A	27	-40.339	91.174	-14.967	1.00	32.59	T8
	ATOM	8204	CG	TRP	A	27	-41.695	91.724	-15.156	1.00	29.33	T8
	ATOM	8205	CD2	TRP	A	27	-42.715	91.212	-16.022	1.00	35.97	T8
40	ATOM	8206	CE2	TRP	A	27	-43.863	92.012	-15.837	1.00	30.81	T8
	ATOM	8207	CE3	TRP	A	27	-42.773	90.154	-16.932	1.00	31.10	T8
	ATOM	8208	CD1	TRP	A	27	-42.244	92.787	-14.503	1.00	33.53	T8
	ATOM	8209	NE1	TRP	A	27	-43.550	92.967	-14.904	1.00	29.41	T8
	ATOM	8210	CZ2	TRP	A	27	-45.052	91.785	-16.529	1.00	33.00	T8
45	ATOM	8211	CZ3	TRP	A	27	-43.958	89.932	-17.620	1.00	32.60	T8
	ATOM	8212	CH2	TRP	A	27	-45.077	90.743	-17.412	1.00	29.54	T8
	ATOM	8213	C	TRP	A	27	-38.920	89.147	-14.704	1.00	32.83	T8
	ATOM	8214	O	TRP	A	27	-38.249	88.840	-13.720	1.00	34.81	T8
	ATOM	8215	N	LEU	A	28	-38.483	89.037	-15.949	1.00	37.51	T8
50	ATOM	8216	CA	LEU	A	28	-37.159	88.553	-16.287	1.00	30.63	T8
	ATOM	8217	CB	LEU	A	28	-37.198	87.052	-16.494	1.00	23.73	T8
	ATOM	8218	CG	LEU	A	28	-35.850	86.359	-16.395	1.00	31.21	T8
	ATOM	8219	CD1	LEU	A	28	-35.300	86.538	-14.979	1.00	27.70	T8
	ATOM	8220	CD2	LEU	A	28	-36.025	84.885	-16.736	1.00	24.39	T8
55	ATOM	8221	C	LEU	A	28	-36.790	89.246	-17.591	1.00	33.08	T8
	ATOM	8222	O	LEU	A	28	-37.564	89.225	-18.548	1.00	36.73	T8
	ATOM	8223	N	LEU	A	29	-35.618	89.867	-17.635	1.00	23.14	T8
	ATOM	8224	CA	LEU	A	29	-35.211	90.584	-18.831	1.00	29.98	T8
	ATOM	8225	CB	LEU	A	29	-33.804	91.141	-18.672	1.00	23.96	T8
60	ATOM	8226	CG	LEU	A	29	-33.330	91.887	-19.911	1.00	26.96	T8
	ATOM	8227	CD1	LEU	A	29	-34.005	93.244	-19.982	1.00	30.20	T8
	ATOM	8228	CD2	LEU	A	29	-31.838	92.034	-19.867	1.00	30.30	T8
	ATOM	8229	C	LEU	A	29	-35.249	89.728	-20.079	1.00	29.72	T8
	ATOM	8230	O	LEU	A	29	-34.663	88.651	-20.120	1.00	34.36	T8
65	ATOM	8231	N	SER	A	30	-35.961	90.207	-21.090	1.00	22.44	T8
	ATOM	8232	CA	SER	A	30	-36.029	89.519	-22.365	1.00	31.21	T8

	ATOM	8233	CB	SER	A	30	-37.324	89.851	-23.093	1.00	20.09	T8
	ATOM	8234	OG	SER	A	30	-37.316	89.263	-24.378	1.00	28.80	T8
	ATOM	8235	C	SER	A	30	-34.846	90.076	-23.147	1.00	30.94	T8
	ATOM	8236	O	SER	A	30	-33.997	89.339	-23.625	1.00	26.57	T8
5	ATOM	8237	N	PHE	A	31	-34.795	91.396	-23.262	1.00	26.96	T8
	ATOM	8238	CA	PHE	A	31	-33.706	92.058	-23.951	1.00	34.17	T8
	ATOM	8239	CB	PHE	A	31	-33.794	91.812	-25.460	1.00	33.47	T8
	ATOM	8240	CG	PHE	A	31	-34.668	92.791	-26.194	1.00	30.79	T8
	ATOM	8241	CD1	PHE	A	31	-34.171	94.024	-26.594	1.00	34.35	T8
10	ATOM	8242	CD2	PHE	A	31	-35.991	92.487	-26.472	1.00	24.60	T8
	ATOM	8243	CE1	PHE	A	31	-34.978	94.937	-27.257	1.00	35.01	T8
	ATOM	8244	CE2	PHE	A	31	-36.804	93.396	-27.135	1.00	37.12	T8
	ATOM	8245	CZ	PHE	A	31	-36.295	94.624	-27.527	1.00	28.33	T8
	ATOM	8246	C	PHE	A	31	-33.777	93.545	-23.644	1.00	26.77	T8
15	ATOM	8247	O	PHE	A	31	-34.843	94.076	-23.343	1.00	39.47	T8
	ATOM	8248	N	LYS	A	32	-32.632	94.210	-23.707	1.00	34.67	T8
	ATOM	8249	CA	LYS	A	32	-32.558	95.639	-23.450	1.00	29.72	T8
	ATOM	8250	CB	LYS	A	32	-31.962	95.899	-22.075	1.00	36.12	T8
	ATOM	8251	CG	LYS	A	32	-31.605	97.343	-21.866	1.00	30.97	T8
20	ATOM	8252	CD	LYS	A	32	-30.814	97.560	-20.598	1.00	30.14	T8
	ATOM	8253	CE	LYS	A	32	-30.251	98.970	-20.596	1.00	33.17	T8
	ATOM	8254	NZ	LYS	A	32	-29.599	99.303	-19.309	1.00	29.03	T8
	ATOM	8255	C	LYS	A	32	-31.691	96.284	-24.525	1.00	33.36	T8
	ATOM	8256	O	LYS	A	32	-30.559	95.874	-24.757	1.00	33.08	T8
25	ATOM	8257	N	ARG	A	33	-32.224	97.301	-25.178	1.00	33.07	T8
	ATOM	8258	CA	ARG	A	33	-31.502	97.973	-26.243	1.00	29.42	T8
	ATOM	8259	CB	ARG	A	33	-32.180	97.641	-27.578	1.00	26.86	T8
	ATOM	8260	CG	ARG	A	33	-31.681	98.377	-28.800	1.00	30.50	T8
	ATOM	8261	CD	ARG	A	33	-32.083	97.601	-30.048	1.00	38.19	T8
30	ATOM	8262	NE	ARG	A	33	-31.717	98.279	-31.287	1.00	30.26	T8
	ATOM	8263	CZ	ARG	A	33	-32.429	99.250	-31.840	1.00	31.78	T8
	ATOM	8264	NH1	ARG	A	33	-33.550	99.650	-31.265	1.00	28.80	T8
	ATOM	8265	NH2	ARG	A	33	-32.011	99.830	-32.953	1.00	29.67	T8
	ATOM	8266	C	ARG	A	33	-31.483	99.471	-26.006	1.00	31.64	T8
35	ATOM	8267	O	ARG	A	33	-32.531	100.098	-25.877	1.00	32.28	T8
	ATOM	8268	N	GLY	A	34	-30.288	100.041	-25.936	1.00	32.79	T8
	ATOM	8269	CA	GLY	A	34	-30.179	101.468	-25.720	1.00	25.85	T8
	ATOM	8270	C	GLY	A	34	-30.014	101.845	-24.265	1.00	36.48	T8
	ATOM	8271	O	GLY	A	34	-29.718	101.002	-23.420	1.00	32.21	T8
40	ATOM	8272	N	SER	A	35	-30.237	103.119	-23.969	1.00	32.49	T8
	ATOM	8273	CA	SER	A	35	-30.072	103.619	-22.615	1.00	28.04	T8
	ATOM	8274	CB	SER	A	35	-28.965	104.667	-22.610	1.00	34.34	T8
	ATOM	8275	OG	SER	A	35	-29.256	105.698	-23.548	1.00	24.26	T8
	ATOM	8276	C	SER	A	35	-31.309	104.228	-21.967	1.00	28.54	T8
45	ATOM	8277	O	SER	A	35	-31.368	104.332	-20.740	1.00	26.63	T8
	ATOM	8278	N	ALA	A	36	-32.289	104.631	-22.772	1.00	26.85	T8
	ATOM	8279	CA	ALA	A	36	-33.491	105.267	-22.241	1.00	34.30	T8
	ATOM	8280	CB	ALA	A	36	-34.344	105.761	-23.379	1.00	31.15	T8
	ATOM	8281	C	ALA	A	36	-34.350	104.456	-21.278	1.00	27.74	T8
50	ATOM	8282	O	ALA	A	36	-35.177	105.028	-20.573	1.00	27.20	T8
	ATOM	8283	N	LEU	A	37	-34.167	103.137	-21.240	1.00	37.16	T8
	ATOM	8284	CA	LEU	A	37	-34.959	102.289	-20.353	1.00	31.19	T8
	ATOM	8285	CB	LEU	A	37	-36.049	101.595	-21.161	1.00	30.08	T8
	ATOM	8286	CG	LEU	A	37	-37.021	102.548	-21.857	1.00	35.31	T8
55	ATOM	8287	CD1	LEU	A	37	-37.757	101.838	-22.965	1.00	30.45	T8
	ATOM	8288	CD2	LEU	A	37	-37.985	103.111	-20.838	1.00	28.94	T8
	ATOM	8289	C	LEU	A	37	-34.129	101.252	-19.599	1.00	34.16	T8
	ATOM	8290	O	LEU	A	37	-33.143	100.740	-20.114	1.00	31.53	T8
	ATOM	8291	N	GLU	A	38	-34.546	100.941	-18.376	1.00	34.28	T8
60	ATOM	8292	CA	GLU	A	38	-33.848	99.980	-17.520	1.00	31.19	T8
	ATOM	8293	CB	GLU	A	38	-32.891	100.709	-16.590	1.00	27.23	T8
	ATOM	8294	CG	GLU	A	38	-31.555	101.085	-17.186	1.00	31.70	T8
	ATOM	8295	CD	GLU	A	38	-30.769	102.022	-16.271	1.00	27.70	T8
	ATOM	8296	OE1	GLU	A	38	-30.858	101.868	-15.024	1.00	36.58	T8
65	ATOM	8297	OE2	GLU	A	38	-30.057	102.908	-16.807	1.00	29.69	T8
	ATOM	8298	C	GLU	A	38	-34.796	99.179	-16.642	1.00	33.94	T8

	ATOM	8299	O	GLU	A	38	-35.945	99.565	-16.458	1.00	31.35	T8
	ATOM	8300	N	GLU	A	39	-34.310	98.064	-16.096	1.00	30.98	T8
	ATOM	8301	CA	GLU	A	39	-35.123	97.250	-15.190	1.00	29.65	T8
	ATOM	8302	CB	GLU	A	39	-34.707	95.794	-15.170	1.00	39.99	T8
5	ATOM	8303	CG	GLU	A	39	-34.403	95.185	-16.479	1.00	31.03	T8
	ATOM	8304	CD	GLU	A	39	-33.376	94.073	-16.320	1.00	28.48	T8
	ATOM	8305	OE1	GLU	A	39	-32.160	94.352	-16.539	1.00	30.38	T8
	ATOM	8306	OE2	GLU	A	39	-33.787	92.935	-15.953	1.00	27.70	T8
	ATOM	8307	C	GLU	A	39	-34.813	97.772	-13.805	1.00	28.33	T8
10	ATOM	8308	O	GLU	A	39	-33.677	98.173	-13.522	1.00	22.09	T8
	ATOM	8309	N	LYS	A	40	-35.803	97.755	-12.929	1.00	31.54	T8
	ATOM	8310	CA	LYS	A	40	-35.567	98.211	-11.579	1.00	26.76	T8
	ATOM	8311	CB	LYS	A	40	-35.551	99.738	-11.504	1.00	31.36	T8
	ATOM	8312	CG	LYS	A	40	-35.251	100.255	-10.097	1.00	31.72	T8
15	ATOM	8313	CD	LYS	A	40	-35.600	101.722	-9.959	1.00	31.17	T8
	ATOM	8314	CE	LYS	A	40	-35.501	102.172	-8.509	1.00	37.17	T8
	ATOM	8315	NZ	LYS	A	40	-35.999	103.575	-8.365	1.00	34.05	T8
	ATOM	8316	C	LYS	A	40	-36.635	97.674	-10.666	1.00	34.28	T8
	ATOM	8317	O	LYS	A	40	-37.766	98.157	-10.665	1.00	28.12	T8
20	ATOM	8318	N	GLU	A	41	-36.271	96.654	-9.902	1.00	36.08	T8
	ATOM	8319	CA	GLU	A	41	-37.193	96.051	-8.958	1.00	28.70	T8
	ATOM	8320	CB	GLU	A	41	-37.381	97.002	-7.775	1.00	30.02	T8
	ATOM	8321	CG	GLU	A	41	-36.047	97.459	-7.206	1.00	29.35	T8
	ATOM	8322	CD	GLU	A	41	-36.164	98.668	-6.308	1.00	28.90	T8
25	ATOM	8323	OE1	GLU	A	41	-36.762	99.693	-6.732	1.00	39.03	T8
	ATOM	8324	OE2	GLU	A	41	-35.644	98.600	-5.175	1.00	33.83	T8
	ATOM	8325	C	GLU	A	41	-38.529	95.712	-9.605	1.00	34.39	T8
	ATOM	8326	O	GLU	A	41	-39.580	96.206	-9.200	1.00	27.86	T8
	ATOM	8327	N	ASN	A	42	-38.463	94.877	-10.632	1.00	35.75	T8
30	ATOM	8328	CA	ASN	A	42	-39.642	94.419	-11.342	1.00	31.01	T8
	ATOM	8329	CB	ASN	A	42	-40.541	93.656	-10.391	1.00	28.70	T8
	ATOM	8330	CG	ASN	A	42	-41.086	92.409	-11.012	1.00	28.37	T8
	ATOM	8331	OD1	ASN	A	42	-42.291	92.186	-11.024	1.00	24.78	T8
	ATOM	8332	ND2	ASN	A	42	-40.199	91.574	-11.537	1.00	29.80	T8
35	ATOM	8333	C	ASN	A	42	-40.450	95.493	-12.038	1.00	33.81	T8
	ATOM	8334	O	ASN	A	42	-41.620	95.288	-12.341	1.00	30.84	T8
	ATOM	8335	N	LYS	A	43	-39.825	96.635	-12.292	1.00	34.17	T8
	ATOM	8336	CA	LYS	A	43	-40.490	97.736	-12.973	1.00	31.12	T8
	ATOM	8337	CB	LYS	A	43	-40.826	98.850	-11.987	1.00	37.63	T8
40	ATOM	8338	CG	LYS	A	43	-41.899	98.490	-10.996	1.00	27.02	T8
	ATOM	8339	CD	LYS	A	43	-42.136	99.607	-10.001	1.00	37.12	T8
	ATOM	8340	CE	LYS	A	43	-40.973	99.720	-9.033	1.00	28.67	T8
	ATOM	8341	NZ	LYS	A	43	-41.188	100.792	-8.011	1.00	23.70	T8
	ATOM	8342	C	LYS	A	43	-39.567	98.274	-14.047	1.00	24.85	T8
45	ATOM	8343	O	LYS	A	43	-38.372	98.020	-14.016	1.00	26.74	T8
	ATOM	8344	N	ILE	A	44	-40.118	99.003	-15.005	1.00	26.70	T8
	ATOM	8345	CA	ILE	A	44	-39.303	99.574	-16.054	1.00	26.86	T8
	ATOM	8346	CB	ILE	A	44	-40.018	99.534	-17.398	1.00	33.21	T8
	ATOM	8347	CG2	ILE	A	44	-39.144	100.154	-18.466	1.00	28.03	T8
50	ATOM	8348	CG1	ILE	A	44	-40.340	98.088	-17.764	1.00	24.38	T8
	ATOM	8349	CD1	ILE	A	44	-41.074	97.935	-19.080	1.00	34.15	T8
	ATOM	8350	C	ILE	A	44	-39.030	101.015	-15.678	1.00	35.84	T8
	ATOM	8351	O	ILE	A	44	-39.953	101.786	-15.442	1.00	32.85	T8
	ATOM	8352	N	LEU	A	45	-37.757	101.373	-15.619	1.00	34.60	T8
55	ATOM	8353	CA	LEU	A	45	-37.357	102.723	-15.256	1.00	26.91	T8
	ATOM	8354	CB	LEU	A	45	-36.145	102.666	-14.337	1.00	32.26	T8
	ATOM	8355	CG	LEU	A	45	-35.589	104.035	-13.946	1.00	34.53	T8
	ATOM	8356	CD1	LEU	A	45	-36.571	104.749	-13.013	1.00	32.87	T8
	ATOM	8357	CD2	LEU	A	45	-34.242	103.846	-13.281	1.00	26.76	T8
60	ATOM	8358	C	LEU	A	45	-37.026	103.603	-16.453	1.00	32.47	T8
	ATOM	8359	O	LEU	A	45	-36.233	103.223	-17.303	1.00	30.53	T8
	ATOM	8360	N	VAL	A	46	-37.621	104.789	-16.504	1.00	29.39	T8
	ATOM	8361	CA	VAL	A	46	-37.383	105.723	-17.596	1.00	35.59	T8
	ATOM	8362	CB	VAL	A	46	-38.578	106.654	-17.786	1.00	31.10	T8
65	ATOM	8363	CG1	VAL	A	46	-38.306	107.612	-18.916	1.00	25.49	T8
	ATOM	8364	CG2	VAL	A	46	-39.815	105.849	-18.075	1.00	32.00	T8

	ATOM	8365	C	VAL	A	46	-36.145	106.569	-17.324	1.00	30.53	T8
	ATOM	8366	O	VAL	A	46	-36.087	107.296	-16.340	1.00	36.62	T8
	ATOM	8367	N	LYS	A	47	-35.156	106.479	-18.204	1.00	30.42	T8
	ATOM	8368	CA	LYS	A	47	-33.918	107.228	-18.030	1.00	26.64	T8
5	ATOM	8369	CB	LYS	A	47	-32.714	106.322	-18.297	1.00	23.92	T8
	ATOM	8370	CG	LYS	A	47	-32.404	105.357	-17.180	1.00	39.16	T8
	ATOM	8371	CD	LYS	A	47	-32.261	106.109	-15.877	1.00	34.10	T8
	ATOM	8372	CE	LYS	A	47	-31.650	105.253	-14.781	1.00	27.09	T8
	ATOM	8373	NZ	LYS	A	47	-30.183	105.020	-14.977	1.00	32.44	T8
10	ATOM	8374	C	LYS	A	47	-33.804	108.479	-18.896	1.00	31.87	T8
	ATOM	8375	O	LYS	A	47	-32.882	109.269	-18.720	1.00	33.89	T8
	ATOM	8376	N	GLU	A	48	-34.732	108.652	-19.832	1.00	32.04	T8
	ATOM	8377	CA	GLU	A	48	-34.732	109.811	-20.722	1.00	34.87	T8
	ATOM	8378	CB	GLU	A	48	-34.096	109.477	-22.060	1.00	35.08	T8
15	ATOM	8379	CG	GLU	A	48	-32.612	109.265	-22.035	1.00	36.49	T8
	ATOM	8380	CD	GLU	A	48	-32.095	108.794	-23.380	1.00	32.30	T8
	ATOM	8381	OE1	GLU	A	48	-32.560	109.339	-24.419	1.00	23.04	T8
	ATOM	8382	OE2	GLU	A	48	-31.227	107.883	-23.395	1.00	31.73	T8
	ATOM	8383	C	GLU	A	48	-36.164	110.167	-20.987	1.00	27.24	T8
20	ATOM	8384	O	GLU	A	48	-36.928	109.319	-21.440	1.00	37.28	T8
	ATOM	8385	N	THR	A	49	-36.543	111.412	-20.730	1.00	29.35	T8
	ATOM	8386	CA	THR	A	49	-37.925	111.779	-20.973	1.00	30.87	T8
	ATOM	8387	CB	THR	A	49	-38.253	113.142	-20.377	1.00	32.31	T8
	ATOM	8388	OG1	THR	A	49	-38.038	114.148	-21.365	1.00	28.82	T8
25	ATOM	8389	CG2	THR	A	49	-37.366	113.417	-19.169	1.00	28.75	T8
	ATOM	8390	C	THR	A	49	-38.195	111.789	-22.478	1.00	29.42	T8
	ATOM	8391	O	THR	A	49	-37.300	112.037	-23.278	1.00	31.34	T8
	ATOM	8392	N	GLY	A	50	-39.434	111.493	-22.852	1.00	27.55	T8
	ATOM	8393	CA	GLY	A	50	-39.803	111.463	-24.254	1.00	27.14	T8
30	ATOM	8394	C	GLY	A	50	-41.116	110.726	-24.445	1.00	25.33	T8
	ATOM	8395	O	GLY	A	50	-41.848	110.510	-23.486	1.00	29.98	T8
	ATOM	8396	N	TYR	A	51	-41.418	110.347	-25.684	1.00	29.95	T8
	ATOM	8397	CA	TYR	A	51	-42.645	109.622	-25.994	1.00	38.10	T8
	ATOM	8398	CB	TYR	A	51	-43.299	110.191	-27.260	1.00	28.65	T8
35	ATOM	8399	CG	TYR	A	51	-43.965	111.515	-27.012	1.00	30.61	T8
	ATOM	8400	CD1	TYR	A	51	-43.221	112.687	-26.936	1.00	33.85	T8
	ATOM	8401	CE1	TYR	A	51	-43.818	113.900	-26.578	1.00	23.85	T8
	ATOM	8402	CD2	TYR	A	51	-45.325	111.582	-26.736	1.00	35.17	T8
	ATOM	8403	CE2	TYR	A	51	-45.933	112.776	-26.375	1.00	30.77	T8
40	ATOM	8404	CZ	TYR	A	51	-45.178	113.932	-26.292	1.00	35.49	T8
	ATOM	8405	OH	TYR	A	51	-45.779	115.104	-25.878	1.00	29.67	T8
	ATOM	8406	C	TYR	A	51	-42.355	108.138	-26.172	1.00	36.58	T8
	ATOM	8407	O	TYR	A	51	-41.469	107.757	-26.940	1.00	32.73	T8
	ATOM	8408	N	PHE	A	52	-43.102	107.301	-25.458	1.00	27.47	T8
45	ATOM	8409	CA	PHE	A	52	-42.901	105.864	-25.538	1.00	30.60	T8
	ATOM	8410	CB	PHE	A	52	-42.385	105.321	-24.209	1.00	31.88	T8
	ATOM	8411	CG	PHE	A	52	-41.109	105.953	-23.742	1.00	29.60	T8
	ATOM	8412	CD1	PHE	A	52	-41.101	107.244	-23.220	1.00	23.78	T8
	ATOM	8413	CD2	PHE	A	52	-39.914	105.246	-23.802	1.00	34.23	T8
50	ATOM	8414	CE1	PHE	A	52	-39.919	107.822	-22.762	1.00	29.86	T8
	ATOM	8415	CE2	PHE	A	52	-38.735	105.811	-23.350	1.00	25.79	T8
	ATOM	8416	CZ	PHE	A	52	-38.736	107.104	-22.827	1.00	28.93	T8
	ATOM	8417	C	PHE	A	52	-44.144	105.080	-25.902	1.00	28.81	T8
	ATOM	8418	O	PHE	A	52	-45.265	105.468	-25.571	1.00	33.92	T8
55	ATOM	8419	N	PHE	A	53	-43.918	103.971	-26.595	1.00	33.68	T8
	ATOM	8420	CA	PHE	A	53	-44.976	103.054	-26.985	1.00	28.90	T8
	ATOM	8421	CB	PHE	A	53	-44.698	102.446	-28.355	1.00	37.34	T8
	ATOM	8422	CG	PHE	A	53	-45.637	101.336	-28.725	1.00	27.24	T8
	ATOM	8423	CD1	PHE	A	53	-46.975	101.594	-28.980	1.00	34.08	T8
60	ATOM	8424	CD2	PHE	A	53	-45.185	100.030	-28.805	1.00	30.98	T8
	ATOM	8425	CE1	PHE	A	53	-47.841	100.570	-29.309	1.00	37.92	T8
	ATOM	8426	CE2	PHE	A	53	-46.045	99.005	-29.130	1.00	32.72	T8
	ATOM	8427	CZ	PHE	A	53	-47.373	99.274	-29.382	1.00	31.20	T8
	ATOM	8428	C	PHE	A	53	-44.862	101.981	-25.922	1.00	36.49	T8
65	ATOM	8429	O	PHE	A	53	-43.800	101.410	-25.744	1.00	26.95	T8
	ATOM	8430	N	ILE	A	54	-45.945	101.714	-25.210	1.00	25.05	T8

	ATOM	8431	CA	ILE	A	54	-45.916	100.730	-24.144	1.00	30.49	T8
	ATOM	8432	CB	ILE	A	54	-46.221	101.407	-22.805	1.00	25.84	T8
	ATOM	8433	CG2	ILE	A	54	-45.969	100.444	-21.658	1.00	31.03	T8
	ATOM	8434	CG1	ILE	A	54	-45.325	102.634	-22.654	1.00	28.24	T8
5	ATOM	8435	CD1	ILE	A	54	-45.810	103.624	-21.643	1.00	32.51	T8
	ATOM	8436	C	ILE	A	54	-46.922	99.626	-24.391	1.00	33.60	T8
	ATOM	8437	O	ILE	A	54	-48.086	99.894	-24.670	1.00	31.55	T8
	ATOM	8438	N	TYR	A	55	-46.473	98.382	-24.282	1.00	30.87	T8
	ATOM	8439	CA	TYR	A	55	-47.352	97.244	-24.513	1.00	28.30	T8
10	ATOM	8440	CB	TYR	A	55	-47.030	96.604	-25.865	1.00	28.66	T8
	ATOM	8441	CG	TYR	A	55	-45.598	96.165	-26.016	1.00	33.04	T8
	ATOM	8442	CD1	TYR	A	55	-45.218	94.858	-25.751	1.00	37.78	T8
	ATOM	8443	CE1	TYR	A	55	-43.896	94.465	-25.852	1.00	38.41	T8
	ATOM	8444	CD2	TYR	A	55	-44.613	97.071	-26.391	1.00	34.82	T8
15	ATOM	8445	CE2	TYR	A	55	-43.283	96.687	-26.493	1.00	26.93	T8
	ATOM	8446	CZ	TYR	A	55	-42.934	95.386	-26.220	1.00	28.49	T8
	ATOM	8447	OH	TYR	A	55	-41.616	95.018	-26.293	1.00	22.82	T8
	ATOM	8448	C	TYR	A	55	-47.263	96.204	-23.415	1.00	28.37	T8
	ATOM	8449	O	TYR	A	55	-46.422	96.295	-22.534	1.00	28.82	T8
20	ATOM	8450	N	GLY	A	56	-48.147	95.220	-23.462	1.00	33.64	T8
	ATOM	8451	CA	GLY	A	56	-48.126	94.187	-22.452	1.00	30.22	T8
	ATOM	8452	C	GLY	A	56	-49.244	93.195	-22.641	1.00	31.48	T8
	ATOM	8453	O	GLY	A	56	-50.356	93.579	-22.976	1.00	31.44	T8
	ATOM	8454	N	GLN	A	57	-48.933	91.917	-22.440	1.00	32.87	T8
25	ATOM	8455	CA	GLN	A	57	-49.906	90.840	-22.566	1.00	36.09	T8
	ATOM	8456	CB	GLN	A	57	-49.702	90.062	-23.874	1.00	32.53	T8
	ATOM	8457	CG	GLN	A	57	-50.585	88.819	-23.988	1.00	31.22	T8
	ATOM	8458	CD	GLN	A	57	-50.480	88.107	-25.319	1.00	26.33	T8
	ATOM	8459	OE1	GLN	A	57	-50.932	88.611	-26.336	1.00	32.47	T8
30	ATOM	8460	NE2	GLN	A	57	-49.885	86.926	-25.315	1.00	35.24	T8
	ATOM	8461	C	GLN	A	57	-49.794	89.884	-21.385	1.00	29.82	T8
	ATOM	8462	O	GLN	A	57	-48.714	89.677	-20.849	1.00	32.67	T8
	ATOM	8463	N	VAL	A	58	-50.928	89.318	-20.988	1.00	33.33	T8
	ATOM	8464	CA	VAL	A	58	-51.002	88.364	-19.889	1.00	26.44	T8
35	ATOM	8465	CB	VAL	A	58	-51.525	89.034	-18.601	1.00	33.75	T8
	ATOM	8466	CG1	VAL	A	58	-51.835	87.990	-17.551	1.00	28.19	T8
	ATOM	8467	CG2	VAL	A	58	-50.502	89.993	-18.069	1.00	32.27	T8
	ATOM	8468	C	VAL	A	58	-51.969	87.242	-20.272	1.00	31.79	T8
	ATOM	8469	O	VAL	A	58	-52.992	87.501	-20.888	1.00	35.06	T8
40	ATOM	8470	N	LEU	A	59	-51.640	86.000	-19.922	1.00	32.68	T8
	ATOM	8471	CA	LEU	A	59	-52.515	84.869	-20.218	1.00	32.16	T8
	ATOM	8472	CB	LEU	A	59	-51.706	83.660	-20.668	1.00	32.73	T8
	ATOM	8473	CG	LEU	A	59	-52.447	82.630	-21.531	1.00	27.50	T8
	ATOM	8474	CD1	LEU	A	59	-51.670	81.337	-21.528	1.00	32.45	T8
45	ATOM	8475	CD2	LEU	A	59	-53.836	82.373	-21.013	1.00	29.10	T8
	ATOM	8476	C	LEU	A	59	-53.265	84.510	-18.937	1.00	28.02	T8
	ATOM	8477	O	LEU	A	59	-52.660	84.065	-17.963	1.00	33.45	T8
	ATOM	8478	N	TYR	A	60	-54.582	84.696	-18.936	1.00	31.23	T8
	ATOM	8479	CA	TYR	A	60	-55.381	84.399	-17.752	1.00	33.53	T8
50	ATOM	8480	CB	TYR	A	60	-56.527	85.391	-17.635	1.00	32.31	T8
	ATOM	8481	CG	TYR	A	60	-56.037	86.797	-17.506	1.00	35.05	T8
	ATOM	8482	CD1	TYR	A	60	-56.125	87.687	-18.573	1.00	31.44	T8
	ATOM	8483	CE1	TYR	A	60	-55.591	88.966	-18.479	1.00	27.26	T8
	ATOM	8484	CD2	TYR	A	60	-55.410	87.220	-16.338	1.00	25.70	T8
55	ATOM	8485	CE2	TYR	A	60	-54.872	88.482	-16.231	1.00	31.82	T8
	ATOM	8486	CZ	TYR	A	60	-54.963	89.350	-17.302	1.00	24.96	T8
	ATOM	8487	OH	TYR	A	60	-54.399	90.595	-17.194	1.00	30.62	T8
	ATOM	8488	C	TYR	A	60	-55.938	82.998	-17.734	1.00	27.54	T8
	ATOM	8489	O	TYR	A	60	-56.595	82.571	-18.676	1.00	29.18	T8
60	ATOM	8490	N	THR	A	61	-55.676	82.281	-16.651	1.00	33.29	T8
	ATOM	8491	CA	THR	A	61	-56.169	80.921	-16.511	1.00	33.08	T8
	ATOM	8492	CB	THR	A	61	-55.016	79.927	-16.379	1.00	27.99	T8
	ATOM	8493	OG1	THR	A	61	-54.148	80.335	-15.321	1.00	37.02	T8
	ATOM	8494	CG2	THR	A	61	-54.227	79.870	-17.673	1.00	35.46	T8
65	ATOM	8495	C	THR	A	61	-57.039	80.858	-15.281	1.00	30.44	T8
	ATOM	8496	O	THR	A	61	-57.341	79.792	-14.770	1.00	31.93	T8

	ATOM	8497	N	ASP	A	62	-57.439	82.031	-14.816	1.00	31.29	T8
	ATOM	8498	CA	ASP	A	62	-58.294	82.175	-13.641	1.00	30.41	T8
	ATOM	8499	CB	ASP	A	62	-58.015	83.540	-12.994	1.00	28.53	T8
	ATOM	8500	CG	ASP	A	62	-58.596	83.668	-11.606	1.00	36.12	T8
5	ATOM	8501	OD1	ASP	A	62	-57.834	84.072	-10.696	1.00	35.37	T8
	ATOM	8502	OD2	ASP	A	62	-59.803	83.379	-11.433	1.00	28.53	T8
	ATOM	8503	C	ASP	A	62	-59.737	82.086	-14.135	1.00	34.21	T8
	ATOM	8504	O	ASP	A	62	-60.028	82.481	-15.259	1.00	34.08	T8
	ATOM	8505	N	LYS	A	63	-60.644	81.570	-13.315	1.00	33.71	T8
10	ATOM	8506	CA	LYS	A	63	-62.027	81.464	-13.758	1.00	37.34	T8
	ATOM	8507	CB	LYS	A	63	-62.541	80.046	-13.536	1.00	32.28	T8
	ATOM	8508	CG	LYS	A	63	-62.524	79.600	-12.092	1.00	33.29	T8
	ATOM	8509	CD	LYS	A	63	-63.087	78.182	-11.941	1.00	27.84	T8
	ATOM	8510	CE	LYS	A	63	-62.260	77.141	-12.735	1.00	39.17	T8
15	ATOM	8511	NZ	LYS	A	63	-62.776	75.730	-12.626	1.00	33.15	T8
	ATOM	8512	C	LYS	A	63	-62.984	82.445	-13.102	1.00	28.78	T8
	ATOM	8513	O	LYS	A	63	-64.191	82.213	-13.085	1.00	36.06	T8
	ATOM	8514	N	THR	A	64	-62.468	83.555	-12.591	1.00	33.59	T8
	ATOM	8515	CA	THR	A	64	-63.349	84.500	-11.932	1.00	27.58	T8
20	ATOM	8516	CB	THR	A	64	-62.623	85.257	-10.784	1.00	21.70	T8
	ATOM	8517	OG1	THR	A	64	-61.396	85.810	-11.265	1.00	29.28	T8
	ATOM	8518	CG2	THR	A	64	-62.340	84.300	-9.615	1.00	36.96	T8
	ATOM	8519	C	THR	A	64	-64.070	85.504	-12.818	1.00	34.27	T8
	ATOM	8520	O	THR	A	64	-63.909	86.700	-12.659	1.00	30.40	T8
25	ATOM	8521	N	TYR	A	65	-64.878	84.996	-13.740	1.00	24.30	T8
	ATOM	8522	CA	TYR	A	65	-65.698	85.804	-14.650	1.00	32.85	T8
	ATOM	8523	CB	TYR	A	65	-66.968	86.254	-13.919	1.00	31.32	T8
	ATOM	8524	CG	TYR	A	65	-66.964	87.692	-13.469	1.00	20.71	T8
	ATOM	8525	CD1	TYR	A	65	-67.485	88.707	-14.284	1.00	30.99	T8
30	ATOM	8526	CE1	TYR	A	65	-67.474	90.051	-13.874	1.00	22.52	T8
	ATOM	8527	CD2	TYR	A	65	-66.430	88.049	-12.232	1.00	25.56	T8
	ATOM	8528	CE2	TYR	A	65	-66.406	89.387	-11.806	1.00	26.27	T8
	ATOM	8529	CZ	TYR	A	65	-66.929	90.384	-12.627	1.00	30.65	T8
	ATOM	8530	OH	TYR	A	65	-66.908	91.700	-12.195	1.00	25.49	T8
35	ATOM	8531	C	TYR	A	65	-65.087	87.008	-15.371	1.00	27.72	T8
	ATOM	8532	O	TYR	A	65	-65.589	87.416	-16.416	1.00	34.11	T8
	ATOM	8533	N	ALA	A	66	-64.024	87.588	-14.834	1.00	29.13	T8
	ATOM	8534	CA	ALA	A	66	-63.412	88.741	-15.481	1.00	30.31	T8
	ATOM	8535	CB	ALA	A	66	-64.274	89.971	-15.267	1.00	37.60	T8
40	ATOM	8536	C	ALA	A	66	-62.006	88.995	-14.970	1.00	34.36	T8
	ATOM	8537	O	ALA	A	66	-61.807	89.260	-13.796	1.00	31.35	T8
	ATOM	8538	N	MET	A	67	-61.027	88.905	-15.857	1.00	27.06	T8
	ATOM	8539	CA	MET	A	67	-59.639	89.147	-15.489	1.00	27.17	T8
	ATOM	8540	CB	MET	A	67	-58.799	87.892	-15.710	1.00	35.79	T8
45	ATOM	8541	CG	MET	A	67	-59.118	86.763	-14.751	1.00	23.31	T8
	ATOM	8542	SD	MET	A	67	-58.848	87.237	-13.039	1.00	27.10	T8
	ATOM	8543	CE	MET	A	67	-57.052	87.068	-12.929	1.00	29.28	T8
	ATOM	8544	C	MET	A	67	-59.100	90.279	-16.338	1.00	28.06	T8
	ATOM	8545	O	MET	A	67	-59.730	90.688	-17.316	1.00	26.44	T8
50	ATOM	8546	N	GLY	A	68	-57.935	90.790	-15.964	1.00	35.77	T8
	ATOM	8547	CA	GLY	A	68	-57.339	91.877	-16.718	1.00	31.61	T8
	ATOM	8548	C	GLY	A	68	-56.152	92.487	-16.006	1.00	32.96	T8
	ATOM	8549	O	GLY	A	68	-55.899	92.171	-14.846	1.00	31.89	T8
	ATOM	8550	N	HIS	A	69	-55.408	93.346	-16.699	1.00	27.20	T8
55	ATOM	8551	CA	HIS	A	69	-54.253	93.997	-16.088	1.00	29.72	T8
	ATOM	8552	CB	HIS	A	69	-52.940	93.319	-16.505	1.00	33.44	T8
	ATOM	8553	CG	HIS	A	69	-52.744	93.202	-17.986	1.00	29.44	T8
	ATOM	8554	CD2	HIS	A	69	-51.952	93.894	-18.837	1.00	32.88	T8
	ATOM	8555	ND1	HIS	A	69	-53.382	92.252	-18.752	1.00	29.80	T8
60	ATOM	8556	CE1	HIS	A	69	-52.989	92.359	-20.007	1.00	34.17	T8
	ATOM	8557	NE2	HIS	A	69	-52.121	93.348	-20.085	1.00	35.30	T8
	ATOM	8558	C	HIS	A	69	-54.184	95.485	-16.394	1.00	36.79	T8
	ATOM	8559	O	HIS	A	69	-54.896	95.986	-17.258	1.00	22.99	T8
	ATOM	8560	N	LEU	A	70	-53.323	96.182	-15.666	1.00	27.66	T8
65	ATOM	8561	CA	LEU	A	70	-53.148	97.615	-15.827	1.00	35.50	T8
	ATOM	8562	CB	LEU	A	70	-53.534	98.340	-14.541	1.00	29.78	T8

	ATOM	8563	CG	LEU	A	70	-54.748	97.859	-13.766	1.00	31.83	T8
	ATOM	8564	CD1	LEU	A	70	-54.794	98.577	-12.442	1.00	30.73	T8
	ATOM	8565	CD2	LEU	A	70	-56.004	98.106	-14.562	1.00	35.49	T8
	ATOM	8566	C	LEU	A	70	-51.695	97.946	-16.109	1.00	20.24	T8
5	ATOM	8567	O	LEU	A	70	-50.789	97.294	-15.590	1.00	28.43	T8
	ATOM	8568	N	ILE	A	71	-51.473	98.956	-16.937	1.00	34.31	T8
	ATOM	8569	CA	ILE	A	71	-50.119	99.404	-17.207	1.00	25.06	T8
	ATOM	8570	CB	ILE	A	71	-49.855	99.547	-18.710	1.00	35.94	T8
	ATOM	8571	CG2	ILE	A	71	-48.564	100.300	-18.942	1.00	31.55	T8
10	ATOM	8572	CG1	ILE	A	71	-49.765	98.159	-19.348	1.00	25.38	T8
	ATOM	8573	CD1	ILE	A	71	-49.533	98.181	-20.830	1.00	30.29	T8
	ATOM	8574	C	ILE	A	71	-50.074	100.761	-16.525	1.00	33.92	T8
	ATOM	8575	O	ILE	A	71	-50.703	101.711	-16.978	1.00	35.42	T8
	ATOM	8576	N	GLN	A	72	-49.341	100.852	-15.423	1.00	32.14	T8
15	ATOM	8577	CA	GLN	A	72	-49.292	102.097	-14.675	1.00	31.33	T8
	ATOM	8578	CB	GLN	A	72	-49.668	101.813	-13.233	1.00	32.89	T8
	ATOM	8579	CG	GLN	A	72	-50.870	100.919	-13.128	1.00	31.78	T8
	ATOM	8580	CD	GLN	A	72	-51.363	100.790	-11.711	1.00	31.27	T8
	ATOM	8581	OE1	GLN	A	72	-50.603	100.427	-10.807	1.00	31.81	T8
20	ATOM	8582	NE2	GLN	A	72	-52.644	101.090	-11.501	1.00	33.98	T8
	ATOM	8583	C	GLN	A	72	-47.992	102.883	-14.716	1.00	37.57	T8
	ATOM	8584	O	GLN	A	72	-46.915	102.341	-14.967	1.00	29.84	T8
	ATOM	8585	N	ARG	A	73	-48.117	104.176	-14.440	1.00	32.00	T8
	ATOM	8586	CA	ARG	A	73	-46.995	105.094	-14.438	1.00	35.13	T8
25	ATOM	8587	CB	ARG	A	73	-47.219	106.147	-15.519	1.00	27.61	T8
	ATOM	8588	CG	ARG	A	73	-46.128	107.165	-15.617	1.00	30.41	T8
	ATOM	8589	CD	ARG	A	73	-46.640	108.436	-16.217	1.00	28.33	T8
	ATOM	8590	NE	ARG	A	73	-45.589	109.439	-16.282	1.00	31.89	T8
	ATOM	8591	CZ	ARG	A	73	-45.806	110.725	-16.511	1.00	34.71	T8
30	ATOM	8592	NH1	ARG	A	73	-47.042	111.163	-16.693	1.00	29.31	T8
	ATOM	8593	NH2	ARG	A	73	-44.788	111.570	-16.567	1.00	30.77	T8
	ATOM	8594	C	ARG	A	73	-46.846	105.788	-13.080	1.00	27.52	T8
	ATOM	8595	O	ARG	A	73	-47.831	106.267	-12.516	1.00	34.56	T8
	ATOM	8596	N	LYS	A	74	-45.620	105.832	-12.557	1.00	30.75	T8
35	ATOM	8597	CA	LYS	A	74	-45.349	106.499	-11.286	1.00	37.71	T8
	ATOM	8598	CB	LYS	A	74	-44.478	105.632	-10.378	1.00	32.46	T8
	ATOM	8599	CG	LYS	A	74	-45.165	104.384	-9.860	1.00	27.06	T8
	ATOM	8600	CD	LYS	A	74	-44.216	103.492	-9.031	1.00	31.56	T8
	ATOM	8601	CE	LYS	A	74	-44.922	102.201	-8.551	1.00	33.21	T8
40	ATOM	8602	NZ	LYS	A	74	-44.047	101.316	-7.701	1.00	28.16	T8
	ATOM	8603	C	LYS	A	74	-44.607	107.790	-11.582	1.00	28.83	T8
	ATOM	8604	O	LYS	A	74	-43.402	107.778	-11.830	1.00	25.96	T8
	ATOM	8605	N	LYS	A	75	-45.329	108.903	-11.561	1.00	27.58	T8
	ATOM	8606	CA	LYS	A	75	-44.735	110.205	-11.831	1.00	28.54	T8
45	ATOM	8607	CB	LYS	A	75	-45.798	111.300	-11.738	1.00	39.72	T8
	ATOM	8608	CG	LYS	A	75	-46.950	111.173	-12.695	1.00	25.61	T8
	ATOM	8609	CD	LYS	A	75	-47.914	112.341	-12.502	1.00	29.90	T8
	ATOM	8610	CE	LYS	A	75	-49.115	112.239	-13.446	1.00	25.79	T8
	ATOM	8611	NZ	LYS	A	75	-50.174	113.279	-13.229	1.00	34.41	T8
50	ATOM	8612	C	LYS	A	75	-43.621	110.536	-10.845	1.00	25.72	T8
	ATOM	8613	O	LYS	A	75	-43.768	110.312	-9.649	1.00	28.55	T8
	ATOM	8614	N	VAL	A	76	-42.515	111.080	-11.344	1.00	34.30	T8
	ATOM	8615	CA	VAL	A	76	-41.408	111.468	-10.475	1.00	33.73	T8
	ATOM	8616	CB	VAL	A	76	-40.124	111.735	-11.238	1.00	28.92	T8
55	ATOM	8617	CG1	VAL	A	76	-38.968	111.734	-10.285	1.00	26.66	T8
	ATOM	8618	CG2	VAL	A	76	-39.947	110.733	-12.317	1.00	27.55	T8
	ATOM	8619	C	VAL	A	76	-41.778	112.798	-9.858	1.00	30.25	T8
	ATOM	8620	O	VAL	A	76	-41.525	113.047	-8.687	1.00	30.46	T8
	ATOM	8621	N	HIS	A	77	-42.362	113.659	-10.681	1.00	27.25	T8
60	ATOM	8622	CA	HIS	A	77	-42.768	114.979	-10.250	1.00	35.56	T8
	ATOM	8623	CB	HIS	A	77	-42.327	116.010	-11.282	1.00	28.46	T8
	ATOM	8624	CG	HIS	A	77	-40.850	116.026	-11.519	1.00	28.37	T8
	ATOM	8625	CD2	HIS	A	77	-40.117	116.492	-12.555	1.00	35.99	T8
	ATOM	8626	ND1	HIS	A	77	-39.944	115.573	-10.583	1.00	30.59	T8
65	ATOM	8627	CE1	HIS	A	77	-38.716	115.759	-11.030	1.00	27.94	T8
	ATOM	8628	NE2	HIS	A	77	-38.794	116.316	-12.224	1.00	28.95	T8

	ATOM	8629	C	HIS	A	77	-44.273	114.994	-10.083	1.00	25.68	T8
	ATOM	8630	O	HIS	A	77	-44.977	114.308	-10.821	1.00	28.13	T8
	ATOM	8631	N	VAL	A	78	-44.773	115.786	-9.134	1.00	33.07	T8
	ATOM	8632	CA	VAL	A	78	-46.197	115.795	-8.893	1.00	25.20	T8
5	ATOM	8633	CB	VAL	A	78	-46.490	115.078	-7.567	1.00	31.96	T8
	ATOM	8634	CG1	VAL	A	78	-47.979	114.925	-7.367	1.00	22.71	T8
	ATOM	8635	CG2	VAL	A	78	-45.857	113.703	-7.588	1.00	39.19	T8
	ATOM	8636	C	VAL	A	78	-46.992	117.101	-8.981	1.00	33.79	T8
	ATOM	8637	O	VAL	A	78	-47.826	117.235	-9.872	1.00	33.00	T8
10	ATOM	8638	N	PHE	A	79	-46.777	118.059	-8.093	1.00	31.62	T8
	ATOM	8639	CA	PHE	A	79	-47.551	119.319	-8.152	1.00	33.29	T8
	ATOM	8640	CB	PHE	A	79	-47.690	119.864	-9.590	1.00	28.26	T8
	ATOM	8641	CG	PHE	A	79	-46.400	120.003	-10.331	1.00	31.49	T8
	ATOM	8642	CD1	PHE	A	79	-46.009	119.039	-11.261	1.00	25.15	T8
15	ATOM	8643	CD2	PHE	A	79	-45.568	121.086	-10.097	1.00	37.83	T8
	ATOM	8644	CE1	PHE	A	79	-44.809	119.154	-11.942	1.00	27.41	T8
	ATOM	8645	CE2	PHE	A	79	-44.367	121.213	-10.772	1.00	31.70	T8
	ATOM	8646	CZ	PHE	A	79	-43.985	120.244	-11.696	1.00	33.72	T8
	ATOM	8647	C	PHE	A	79	-48.981	119.222	-7.600	1.00	24.51	T8
20	ATOM	8648	O	PHE	A	79	-49.795	118.413	-8.063	1.00	28.10	T8
	ATOM	8649	N	GLY	A	80	-49.287	120.086	-6.635	1.00	33.12	T8
	ATOM	8650	CA	GLY	A	80	-50.614	120.129	-6.039	1.00	31.00	T8
	ATOM	8651	C	GLY	A	80	-51.248	118.807	-5.650	1.00	29.31	T8
	ATOM	8652	O	GLY	A	80	-50.635	117.972	-4.974	1.00	32.24	T8
25	ATOM	8653	N	ASP	A	81	-52.491	118.620	-6.088	1.00	28.96	T8
	ATOM	8654	CA	ASP	A	81	-53.247	117.410	-5.782	1.00	32.27	T8
	ATOM	8655	CB	ASP	A	81	-54.694	117.778	-5.432	1.00	33.27	T8
	ATOM	8656	CG	ASP	A	81	-55.455	118.359	-6.607	1.00	33.08	T8
	ATOM	8657	OD1	ASP	A	81	-54.818	118.788	-7.587	1.00	28.53	T8
30	ATOM	8658	OD2	ASP	A	81	-56.699	118.402	-6.550	1.00	28.22	T8
	ATOM	8659	C	ASP	A	81	-53.229	116.346	-6.880	1.00	34.51	T8
	ATOM	8660	O	ASP	A	81	-54.176	115.580	-7.023	1.00	35.52	T8
	ATOM	8661	N	GLU	A	82	-52.157	116.301	-7.659	1.00	27.76	T8
	ATOM	8662	CA	GLU	A	82	-52.054	115.298	-8.706	1.00	33.25	T8
35	ATOM	8663	CB	GLU	A	82	-50.832	115.537	-9.586	1.00	36.29	T8
	ATOM	8664	CG	GLU	A	82	-50.943	116.580	-10.641	1.00	27.87	T8
	ATOM	8665	CD	GLU	A	82	-50.016	116.265	-11.794	1.00	33.54	T8
	ATOM	8666	OE1	GLU	A	82	-48.864	115.873	-11.532	1.00	38.53	T8
	ATOM	8667	OE2	GLU	A	82	-50.430	116.399	-12.963	1.00	32.13	T8
40	ATOM	8668	C	GLU	A	82	-51.841	113.963	-8.020	1.00	34.57	T8
	ATOM	8669	O	GLU	A	82	-51.272	113.921	-6.940	1.00	33.08	T8
	ATOM	8670	N	LEU	A	83	-52.292	112.876	-8.630	1.00	33.03	T8
	ATOM	8671	CA	LEU	A	83	-52.038	111.567	-8.055	1.00	32.92	T8
	ATOM	8672	CB	LEU	A	83	-53.153	110.577	-8.377	1.00	33.27	T8
45	ATOM	8673	CG	LEU	A	83	-54.509	110.809	-7.724	1.00	29.28	T8
	ATOM	8674	CD1	LEU	A	83	-55.092	112.126	-8.216	1.00	25.63	T8
	ATOM	8675	CD2	LEU	A	83	-55.434	109.665	-8.075	1.00	20.54	T8
	ATOM	8676	C	LEU	A	83	-50.772	111.162	-8.778	1.00	34.82	T8
	ATOM	8677	O	LEU	A	83	-50.699	111.275	-10.003	1.00	27.64	T8
50	ATOM	8678	N	SER	A	84	-49.772	110.706	-8.036	1.00	25.69	T8
	ATOM	8679	CA	SER	A	84	-48.513	110.324	-8.653	1.00	34.01	T8
	ATOM	8680	CB	SER	A	84	-47.392	110.349	-7.620	1.00	33.30	T8
	ATOM	8681	OG	SER	A	84	-47.785	109.691	-6.436	1.00	34.53	T8
	ATOM	8682	C	SER	A	84	-48.553	108.977	-9.341	1.00	25.21	T8
55	ATOM	8683	O	SER	A	84	-47.574	108.572	-9.972	1.00	27.56	T8
	ATOM	8684	N	LEU	A	85	-49.681	108.282	-9.231	1.00	32.65	T8
	ATOM	8685	CA	LEU	A	85	-49.829	106.981	-9.869	1.00	28.36	T8
	ATOM	8686	CB	LEU	A	85	-50.144	105.915	-8.822	1.00	26.69	T8
	ATOM	8687	CG	LEU	A	85	-49.969	104.423	-9.156	1.00	32.21	T8
60	ATOM	8688	CD1	LEU	A	85	-51.021	103.967	-10.133	1.00	26.36	T8
	ATOM	8689	CD2	LEU	A	85	-48.582	104.176	-9.708	1.00	30.66	T8
	ATOM	8690	C	LEU	A	85	-50.950	107.069	-10.890	1.00	27.91	T8
	ATOM	8691	O	LEU	A	85	-52.116	107.154	-10.532	1.00	39.67	T8
	ATOM	8692	N	VAL	A	86	-50.589	107.075	-12.167	1.00	33.25	T8
65	ATOM	8693	CA	VAL	A	86	-51.580	107.153	-13.234	1.00	34.82	T8
	ATOM	8694	CB	VAL	A	86	-51.210	108.214	-14.291	1.00	32.37	T8

	ATOM	8695	CG1	VAL	A	86	-52.213	108.175	-15.436	1.00	29.77	T8
	ATOM	8696	CG2	VAL	A	86	-51.178	109.590	-13.662	1.00	32.80	T8
	ATOM	8697	C	VAL	A	86	-51.621	105.816	-13.935	1.00	27.02	T8
	ATOM	8698	O	VAL	A	86	-50.580	105.203	-14.163	1.00	27.65	T8
5	ATOM	8699	N	THR	A	87	-52.815	105.354	-14.271	1.00	32.87	T8
	ATOM	8700	CA	THR	A	87	-52.915	104.094	-14.971	1.00	33.63	T8
	ATOM	8701	CB	THR	A	87	-53.921	103.135	-14.290	1.00	30.49	T8
	ATOM	8702	OG1	THR	A	87	-55.001	102.856	-15.180	1.00	37.92	T8
	ATOM	8703	CG2	THR	A	87	-54.455	103.742	-13.007	1.00	24.58	T8
10	ATOM	8704	C	THR	A	87	-53.304	104.372	-16.420	1.00	37.49	T8
	ATOM	8705	O	THR	A	87	-54.344	104.954	-16.705	1.00	33.71	T8
	ATOM	8706	N	LEU	A	88	-52.423	103.980	-17.331	1.00	24.03	T8
	ATOM	8707	CA	LEU	A	88	-52.617	104.162	-18.764	1.00	21.76	T8
	ATOM	8708	CB	LEU	A	88	-51.264	104.446	-19.433	1.00	34.59	T8
15	ATOM	8709	CG	LEU	A	88	-50.297	105.576	-19.051	1.00	25.66	T8
	ATOM	8710	CD1	LEU	A	88	-50.422	105.951	-17.617	1.00	33.13	T8
	ATOM	8711	CD2	LEU	A	88	-48.894	105.125	-19.327	1.00	36.67	T8
	ATOM	8712	C	LEU	A	88	-53.137	102.842	-19.310	1.00	34.12	T8
	ATOM	8713	O	LEU	A	88	-52.778	101.763	-18.820	1.00	30.24	T8
20	ATOM	8714	N	PHE	A	89	-54.001	102.890	-20.303	1.00	34.50	T8
	ATOM	8715	CA	PHE	A	89	-54.442	101.629	-20.903	1.00	32.65	T8
	ATOM	8716	CB	PHE	A	89	-53.211	100.896	-21.437	1.00	28.94	T8
	ATOM	8717	CG	PHE	A	89	-52.250	101.809	-22.135	1.00	30.28	T8
	ATOM	8718	CD1	PHE	A	89	-50.880	101.594	-22.056	1.00	22.02	T8
25	ATOM	8719	CD2	PHE	A	89	-52.723	102.963	-22.798	1.00	24.22	T8
	ATOM	8720	CE1	PHE	A	89	-49.987	102.527	-22.617	1.00	25.48	T8
	ATOM	8721	CE2	PHE	A	89	-51.846	103.895	-23.360	1.00	26.99	T8
	ATOM	8722	CZ	PHE	A	89	-50.477	103.682	-23.267	1.00	28.25	T8
	ATOM	8723	C	PHE	A	89	-55.315	100.659	-20.097	1.00	28.48	T8
30	ATOM	8724	O	PHE	A	89	-56.509	100.915	-19.938	1.00	28.32	T8
	ATOM	8725	N	ARG	A	90	-54.767	99.543	-19.616	1.00	24.49	T8
	ATOM	8726	CA	ARG	A	90	-55.621	98.586	-18.888	1.00	31.85	T8
	ATOM	8727	CB	ARG	A	90	-56.349	99.295	-17.748	1.00	32.63	T8
	ATOM	8728	CG	ARG	A	90	-57.660	98.650	-17.357	1.00	37.11	T8
35	ATOM	8729	CD	ARG	A	90	-58.498	99.597	-16.535	1.00	26.41	T8
	ATOM	8730	NE	ARG	A	90	-59.844	99.068	-16.356	1.00	30.03	T8
	ATOM	8731	CZ	ARG	A	90	-60.949	99.811	-16.338	1.00	24.39	T8
	ATOM	8732	NH1	ARG	A	90	-60.873	101.134	-16.483	1.00	34.36	T8
	ATOM	8733	NH2	ARG	A	90	-62.137	99.224	-16.206	1.00	32.49	T8
40	ATOM	8734	C	ARG	A	90	-56.683	97.862	-19.760	1.00	37.37	T8
	ATOM	8735	O	ARG	A	90	-57.465	98.499	-20.468	1.00	27.05	T8
	ATOM	8736	N	CYS	A	91	-56.723	96.532	-19.675	1.00	28.11	T8
	ATOM	8737	CA	CYS	A	91	-57.681	95.728	-20.443	1.00	27.36	T8
	ATOM	8738	CB	CYS	A	91	-56.957	94.990	-21.568	1.00	31.20	T8
45	ATOM	8739	SG	CYS	A	91	-55.583	93.986	-20.988	1.00	28.14	T8
	ATOM	8740	C	CYS	A	91	-58.435	94.722	-19.564	1.00	26.79	T8
	ATOM	8741	O	CYS	A	91	-58.042	94.466	-18.429	1.00	35.72	T8
	ATOM	8742	N	ILE	A	92	-59.513	94.155	-20.100	1.00	28.61	T8
	ATOM	8743	CA	ILE	A	92	-60.345	93.200	-19.360	1.00	27.40	T8
50	ATOM	8744	CB	ILE	A	92	-61.633	93.871	-18.852	1.00	27.72	T8
	ATOM	8745	CG2	ILE	A	92	-62.444	92.899	-18.027	1.00	26.38	T8
	ATOM	8746	CG1	ILE	A	92	-61.284	95.091	-18.007	1.00	22.42	T8
	ATOM	8747	CD1	ILE	A	92	-62.474	96.002	-17.732	1.00	25.37	T8
	ATOM	8748	C	ILE	A	92	-60.776	92.057	-20.257	1.00	30.79	T8
55	ATOM	8749	O	ILE	A	92	-60.858	92.222	-21.460	1.00	35.92	T8
	ATOM	8750	N	GLN	A	93	-61.058	90.901	-19.671	1.00	30.70	T8
	ATOM	8751	CA	GLN	A	93	-61.513	89.746	-20.443	1.00	30.56	T8
	ATOM	8752	CB	GLN	A	93	-60.322	88.931	-20.955	1.00	31.48	T8
	ATOM	8753	CG	GLN	A	93	-59.749	89.409	-22.279	1.00	27.59	T8
60	ATOM	8754	CD	GLN	A	93	-60.266	88.631	-23.475	1.00	28.25	T8
	ATOM	8755	OE1	GLN	A	93	-61.396	88.814	-23.926	1.00	25.45	T8
	ATOM	8756	NE2	GLN	A	93	-59.428	87.746	-23.997	1.00	32.75	T8
	ATOM	8757	C	GLN	A	93	-62.418	88.847	-19.610	1.00	29.48	T8
	ATOM	8758	O	GLN	A	93	-62.016	88.356	-18.558	1.00	31.31	T8
65	ATOM	8759	N	ASN	A	94	-63.648	88.647	-20.073	1.00	33.65	T8
	ATOM	8760	CA	ASN	A	94	-64.569	87.780	-19.361	1.00	30.49	T8

	ATOM	8761	CB	ASN	A	94	-65.930	87.741	-20.064	1.00	31.12	T8
	ATOM	8762	CG	ASN	A	94	-66.799	88.928	-19.721	1.00	33.75	T8
	ATOM	8763	OD1	ASN	A	94	-67.038	89.215	-18.554	1.00	33.65	T8
	ATOM	8764	ND2	ASN	A	94	-67.289	89.616	-20.738	1.00	35.22	T8
5	ATOM	8765	C	ASN	A	94	-63.935	86.393	-19.375	1.00	33.07	T8
	ATOM	8766	O	ASN	A	94	-63.269	86.019	-20.340	1.00	32.82	T8
	ATOM	8767	N	MET	A	95	-64.128	85.633	-18.304	1.00	22.87	T8
	ATOM	8768	CA	MET	A	95	-63.568	84.292	-18.212	1.00	37.89	T8
	ATOM	8769	CB	MET	A	95	-62.724	84.149	-16.943	1.00	42.45	T8
10	ATOM	8770	CG	MET	A	95	-61.545	85.082	-16.873	1.00	21.82	T8
	ATOM	8771	SD	MET	A	95	-60.510	84.989	-18.358	1.00	26.32	T8
	ATOM	8772	CE	MET	A	95	-59.654	83.452	-18.118	1.00	28.76	T8
	ATOM	8773	C	MET	A	95	-64.672	83.246	-18.186	1.00	28.50	T8
	ATOM	8774	O	MET	A	95	-65.791	83.525	-17.744	1.00	27.87	T8
15	ATOM	8775	N	PRO	A	96	-64.377	82.030	-18.681	1.00	30.58	T8
	ATOM	8776	CD	PRO	A	96	-63.149	81.641	-19.379	1.00	24.49	T8
	ATOM	8777	CA	PRO	A	96	-65.331	80.921	-18.711	1.00	27.20	T8
	ATOM	8778	CB	PRO	A	96	-64.706	79.922	-19.678	1.00	26.80	T8
	ATOM	8779	CG	PRO	A	96	-63.695	80.711	-20.415	1.00	30.87	T8
20	ATOM	8780	C	PRO	A	96	-65.336	80.359	-17.309	1.00	23.94	T8
	ATOM	8781	O	PRO	A	96	-64.743	80.940	-16.395	1.00	25.78	T8
	ATOM	8782	N	GLU	A	97	-65.971	79.213	-17.135	1.00	31.27	T8
	ATOM	8783	CA	GLU	A	97	-66.027	78.623	-15.818	1.00	26.74	T8
	ATOM	8784	CB	GLU	A	97	-67.479	78.392	-15.425	1.00	34.17	T8
25	ATOM	8785	CG	GLU	A	97	-67.652	78.133	-13.955	1.00	27.11	T8
	ATOM	8786	CD	GLU	A	97	-68.838	78.893	-13.399	1.00	34.91	T8
	ATOM	8787	OE1	GLU	A	97	-69.972	78.661	-13.897	1.00	26.57	T8
	ATOM	8788	OE2	GLU	A	97	-68.638	79.725	-12.473	1.00	39.62	T8
	ATOM	8789	C	GLU	A	97	-65.274	77.315	-15.805	1.00	30.58	T8
30	ATOM	8790	O	GLU	A	97	-64.820	76.850	-14.759	1.00	31.80	T8
	ATOM	8791	N	THR	A	98	-65.124	76.738	-16.988	1.00	32.49	T8
	ATOM	8792	CA	THR	A	98	-64.457	75.460	-17.119	1.00	29.90	T8
	ATOM	8793	CB	THR	A	98	-65.123	74.642	-18.203	1.00	27.25	T8
	ATOM	8794	OG1	THR	A	98	-64.976	75.332	-19.452	1.00	25.51	T8
35	ATOM	8795	CG2	THR	A	98	-66.601	74.466	-17.894	1.00	32.97	T8
	ATOM	8796	C	THR	A	98	-62.959	75.511	-17.418	1.00	32.78	T8
	ATOM	8797	O	THR	A	98	-62.142	75.210	-16.543	1.00	32.73	T8
	ATOM	8798	N	LEU	A	99	-62.583	75.890	-18.634	1.00	33.85	T8
	ATOM	8799	CA	LEU	A	99	-61.169	75.892	-18.951	1.00	30.00	T8
40	ATOM	8800	CB	LEU	A	99	-60.917	74.924	-20.100	1.00	27.80	T8
	ATOM	8801	CG	LEU	A	99	-61.276	73.485	-19.728	1.00	31.86	T8
	ATOM	8802	CD1	LEU	A	99	-61.171	72.571	-20.934	1.00	34.67	T8
	ATOM	8803	CD2	LEU	A	99	-60.347	73.029	-18.625	1.00	25.82	T8
	ATOM	8804	C	LEU	A	99	-60.586	77.257	-19.268	1.00	26.98	T8
45	ATOM	8805	O	LEU	A	99	-60.259	77.557	-20.422	1.00	31.72	T8
	ATOM	8806	N	PRO	A	100	-60.420	78.100	-18.236	1.00	35.04	T8
	ATOM	8807	CD	PRO	A	100	-60.669	77.798	-16.818	1.00	24.63	T8
	ATOM	8808	CA	PRO	A	100	-59.875	79.448	-18.380	1.00	29.02	T8
	ATOM	8809	CB	PRO	A	100	-59.591	79.847	-16.939	1.00	31.34	T8
50	ATOM	8810	CG	PRO	A	100	-60.695	79.177	-16.207	1.00	36.67	T8
	ATOM	8811	C	PRO	A	100	-58.630	79.503	-19.258	1.00	31.77	T8
	ATOM	8812	O	PRO	A	100	-57.637	78.828	-18.995	1.00	30.95	T8
	ATOM	8813	N	ASN	A	101	-58.704	80.314	-20.303	1.00	30.74	T8
	ATOM	8814	CA	ASN	A	101	-57.600	80.489	-21.220	1.00	28.39	T8
55	ATOM	8815	CB	ASN	A	101	-57.489	79.297	-22.140	1.00	32.36	T8
	ATOM	8816	CG	ASN	A	101	-56.651	78.215	-21.563	1.00	29.66	T8
	ATOM	8817	OD1	ASN	A	101	-55.436	78.360	-21.432	1.00	30.06	T8
	ATOM	8818	ND2	ASN	A	101	-57.286	77.111	-21.198	1.00	31.62	T8
	ATOM	8819	C	ASN	A	101	-57.828	81.711	-22.063	1.00	32.23	T8
60	ATOM	8820	O	ASN	A	101	-58.303	81.590	-23.188	1.00	31.82	T8
	ATOM	8821	N	ASN	A	102	-57.488	82.892	-21.559	1.00	31.19	T8
	ATOM	8822	CA	ASN	A	102	-57.736	84.040	-22.394	1.00	31.84	T8
	ATOM	8823	CB	ASN	A	102	-58.856	84.892	-21.795	1.00	28.98	T8
	ATOM	8824	CG	ASN	A	102	-60.238	84.373	-22.187	1.00	28.15	T8
65	ATOM	8825	OD1	ASN	A	102	-60.483	84.052	-23.351	1.00	29.24	T8
	ATOM	8826	ND2	ASN	A	102	-61.141	84.289	-21.218	1.00	32.25	T8

	ATOM	8827	C	ASN	A	102	-56.606	84.913	-22.921	1.00	29.30	T8
	ATOM	8828	O	ASN	A	102	-56.608	85.217	-24.114	1.00	33.81	T8
	ATOM	8829	N	SER	A	103	-55.638	85.326	-22.119	1.00	32.74	T8
	ATOM	8830	CA	SER	A	103	-54.599	86.159	-22.730	1.00	32.88	T8
5	ATOM	8831	CB	SER	A	103	-53.873	85.368	-23.834	1.00	29.08	T8
	ATOM	8832	OG	SER	A	103	-53.738	86.119	-25.030	1.00	27.36	T8
	ATOM	8833	C	SER	A	103	-55.189	87.455	-23.325	1.00	24.47	T8
	ATOM	8834	O	SER	A	103	-56.160	87.440	-24.086	1.00	30.94	T8
	ATOM	8835	N	CYS	A	104	-54.587	88.584	-22.986	1.00	28.92	T8
10	ATOM	8836	CA	CYS	A	104	-55.091	89.841	-23.480	1.00	32.96	T8
	ATOM	8837	CB	CYS	A	104	-56.096	90.395	-22.483	1.00	27.06	T8
	ATOM	8838	SG	CYS	A	104	-57.050	91.763	-23.088	1.00	26.25	T8
	ATOM	8839	C	CYS	A	104	-53.941	90.806	-23.676	1.00	26.77	T8
	ATOM	8840	O	CYS	A	104	-53.139	91.011	-22.778	1.00	33.55	T8
15	ATOM	8841	N	TYR	A	105	-53.867	91.385	-24.868	1.00	28.90	T8
	ATOM	8842	CA	TYR	A	105	-52.820	92.334	-25.218	1.00	35.20	T8
	ATOM	8843	CB	TYR	A	105	-52.226	91.972	-26.586	1.00	31.92	T8
	ATOM	8844	CG	TYR	A	105	-51.193	92.945	-27.113	1.00	32.71	T8
	ATOM	8845	CD1	TYR	A	105	-49.841	92.737	-26.899	1.00	33.70	T8
20	ATOM	8846	CE1	TYR	A	105	-48.892	93.625	-27.392	1.00	34.48	T8
	ATOM	8847	CD2	TYR	A	105	-51.572	94.073	-27.834	1.00	28.08	T8
	ATOM	8848	CE2	TYR	A	105	-50.628	94.966	-28.323	1.00	30.85	T8
	ATOM	8849	CZ	TYR	A	105	-49.296	94.731	-28.100	1.00	40.35	T8
	ATOM	8850	OH	TYR	A	105	-48.360	95.590	-28.599	1.00	28.19	T8
25	ATOM	8851	C	TYR	A	105	-53.411	93.731	-25.289	1.00	25.48	T8
	ATOM	8852	O	TYR	A	105	-54.560	93.907	-25.669	1.00	29.72	T8
	ATOM	8853	N	SER	A	106	-52.622	94.725	-24.922	1.00	32.51	T8
	ATOM	8854	CA	SER	A	106	-53.068	96.104	-24.989	1.00	33.20	T8
	ATOM	8855	CB	SER	A	106	-53.950	96.442	-23.793	1.00	26.53	T8
30	ATOM	8856	OG	SER	A	106	-54.592	97.695	-23.958	1.00	32.20	T8
	ATOM	8857	C	SER	A	106	-51.817	96.961	-24.996	1.00	32.74	T8
	ATOM	8858	O	SER	A	106	-50.825	96.618	-24.350	1.00	29.66	T8
	ATOM	8859	N	ALA	A	107	-51.859	98.062	-25.741	1.00	28.32	T8
	ATOM	8860	CA	ALA	A	107	-50.718	98.958	-25.841	1.00	27.65	T8
35	ATOM	8861	CB	ALA	A	107	-49.748	98.433	-26.882	1.00	28.36	T8
	ATOM	8862	C	ALA	A	107	-51.157	100.369	-26.201	1.00	35.13	T8
	ATOM	8863	O	ALA	A	107	-52.283	100.586	-26.626	1.00	26.42	T8
	ATOM	8864	N	GLY	A	108	-50.256	101.325	-26.022	1.00	30.67	T8
	ATOM	8865	CA	GLY	A	108	-50.559	102.704	-26.342	1.00	25.77	T8
40	ATOM	8866	C	GLY	A	108	-49.329	103.582	-26.227	1.00	28.28	T8
	ATOM	8867	O	GLY	A	108	-48.259	103.102	-25.880	1.00	32.41	T8
	ATOM	8868	N	ILE	A	109	-49.477	104.869	-26.517	1.00	30.31	T8
	ATOM	8869	CA	ILE	A	109	-48.366	105.804	-26.438	1.00	38.55	T8
	ATOM	8870	CB	ILE	A	109	-48.235	106.622	-27.740	1.00	30.50	T8
45	ATOM	8871	CG2	ILE	A	109	-47.090	107.609	-27.627	1.00	25.53	T8
	ATOM	8872	CG1	ILE	A	109	-48.003	105.689	-28.921	1.00	30.28	T8
	ATOM	8873	CD1	ILE	A	109	-48.081	106.376	-30.240	1.00	36.24	T8
	ATOM	8874	C	ILE	A	109	-48.595	106.768	-25.284	1.00	25.24	T8
	ATOM	8875	O	ILE	A	109	-49.722	107.200	-25.040	1.00	29.39	T8
50	ATOM	8876	N	ALA	A	110	-47.524	107.105	-24.576	1.00	28.55	T8
	ATOM	8877	CA	ALA	A	110	-47.624	108.034	-23.460	1.00	28.95	T8
	ATOM	8878	CB	ALA	A	110	-47.879	107.278	-22.183	1.00	32.06	T8
	ATOM	8879	C	ALA	A	110	-46.337	108.823	-23.339	1.00	27.51	T8
	ATOM	8880	O	ALA	A	110	-45.287	108.370	-23.787	1.00	29.07	T8
55	ATOM	8881	N	LYS	A	111	-46.411	110.010	-22.751	1.00	31.51	T8
	ATOM	8882	CA	LYS	A	111	-45.205	110.800	-22.571	1.00	30.55	T8
	ATOM	8883	CB	LYS	A	111	-45.488	112.286	-22.727	1.00	37.28	T8
	ATOM	8884	CG	LYS	A	111	-44.222	113.120	-22.600	1.00	29.09	T8
	ATOM	8885	CD	LYS	A	111	-44.492	114.602	-22.801	1.00	33.79	T8
60	ATOM	8886	CE	LYS	A	111	-43.205	115.426	-22.692	1.00	28.08	T8
	ATOM	8887	NZ	LYS	A	111	-43.446	116.902	-22.925	1.00	32.49	T8
	ATOM	8888	C	LYS	A	111	-44.699	110.521	-21.168	1.00	27.65	T8
	ATOM	8889	O	LYS	A	111	-45.473	110.555	-20.222	1.00	29.75	T8
	ATOM	8890	N	LEU	A	112	-43.408	110.239	-21.041	1.00	29.31	T8
65	ATOM	8891	CA	LEU	A	112	-42.795	109.928	-19.753	1.00	24.38	T8
	ATOM	8892	CB	LEU	A	112	-42.370	108.459	-19.736	1.00	33.64	T8

	ATOM	8893	CG	LEU	A	112	-43.403	107.430	-20.181	1.00	29.64	T8
	ATOM	8894	CD1	LEU	A	112	-42.740	106.088	-20.398	1.00	35.23	T8
	ATOM	8895	CD2	LEU	A	112	-44.492	107.340	-19.136	1.00	35.88	T8
	ATOM	8896	C	LEU	A	112	-41.563	110.798	-19.508	1.00	31.95	T8
5	ATOM	8897	O	LEU	A	112	-41.007	111.367	-20.446	1.00	33.36	T8
	ATOM	8898	N	GLU	A	113	-41.133	110.885	-18.251	1.00	28.86	T8
	ATOM	8899	CA	GLU	A	113	-39.958	111.676	-17.893	1.00	32.21	T8
	ATOM	8900	CB	GLU	A	113	-40.333	112.874	-17.043	1.00	31.61	T8
	ATOM	8901	CG	GLU	A	113	-41.582	113.577	-17.424	1.00	37.22	T8
10	ATOM	8902	CD	GLU	A	113	-41.621	114.976	-16.838	1.00	26.95	T8
	ATOM	8903	OE1	GLU	A	113	-41.287	115.141	-15.627	1.00	30.22	T8
	ATOM	8904	OE2	GLU	A	113	-41.990	115.915	-17.598	1.00	30.18	T8
	ATOM	8905	C	GLU	A	113	-38.943	110.890	-17.086	1.00	21.76	T8
	ATOM	8906	O	GLU	A	113	-39.285	109.898	-16.444	1.00	32.39	T8
15	ATOM	8907	N	GLU	A	114	-37.698	111.359	-17.103	1.00	35.61	T8
	ATOM	8908	CA	GLU	A	114	-36.626	110.738	-16.339	1.00	24.97	T8
	ATOM	8909	CB	GLU	A	114	-35.465	111.691	-16.162	1.00	35.33	T8
	ATOM	8910	CG	GLU	A	114	-34.424	111.676	-17.222	1.00	32.62	T8
	ATOM	8911	CD	GLU	A	114	-33.143	112.305	-16.709	1.00	34.72	T8
20	ATOM	8912	OE1	GLU	A	114	-32.549	111.734	-15.764	1.00	34.12	T8
	ATOM	8913	OE2	GLU	A	114	-32.741	113.369	-17.235	1.00	38.95	T8
	ATOM	8914	C	GLU	A	114	-37.115	110.428	-14.942	1.00	33.37	T8
	ATOM	8915	O	GLU	A	114	-37.577	111.325	-14.239	1.00	28.88	T8
	ATOM	8916	N	GLY	A	115	-37.000	109.175	-14.525	1.00	28.42	T8
25	ATOM	8917	CA	GLY	A	115	-37.418	108.832	-13.181	1.00	29.19	T8
	ATOM	8918	C	GLY	A	115	-38.776	108.184	-13.097	1.00	31.03	T8
	ATOM	8919	O	GLY	A	115	-39.105	107.587	-12.076	1.00	36.24	T8
	ATOM	8920	N	ASP	A	116	-39.590	108.319	-14.136	1.00	34.66	T8
	ATOM	8921	CA	ASP	A	116	-40.895	107.686	-14.097	1.00	31.41	T8
30	ATOM	8922	CB	ASP	A	116	-41.743	108.079	-15.311	1.00	29.04	T8
	ATOM	8923	CG	ASP	A	116	-42.224	109.512	-15.260	1.00	27.39	T8
	ATOM	8924	OD1	ASP	A	116	-42.347	110.064	-14.145	1.00	36.41	T8
	ATOM	8925	OD2	ASP	A	116	-42.502	110.080	-16.340	1.00	32.89	T8
	ATOM	8926	C	ASP	A	116	-40.668	106.176	-14.112	1.00	25.07	T8
35	ATOM	8927	O	ASP	A	116	-39.650	105.694	-14.605	1.00	24.05	T8
	ATOM	8928	N	GLU	A	117	-41.609	105.426	-13.561	1.00	28.07	T8
	ATOM	8929	CA	GLU	A	117	-41.486	103.982	-13.562	1.00	32.59	T8
	ATOM	8930	CB	GLU	A	117	-41.269	103.458	-12.147	1.00	36.02	T8
	ATOM	8931	CG	GLU	A	117	-40.099	104.081	-11.435	1.00	34.17	T8
40	ATOM	8932	CD	GLU	A	117	-39.860	103.463	-10.072	1.00	30.17	T8
	ATOM	8933	OE1	GLU	A	117	-40.861	103.131	-9.391	1.00	32.73	T8
	ATOM	8934	OE2	GLU	A	117	-38.673	103.322	-9.679	1.00	29.17	T8
	ATOM	8935	C	GLU	A	117	-42.767	103.401	-14.122	1.00	37.75	T8
	ATOM	8936	O	GLU	A	117	-43.853	103.913	-13.850	1.00	29.34	T8
45	ATOM	8937	N	LEU	A	118	-42.631	102.345	-14.917	1.00	30.38	T8
	ATOM	8938	CA	LEU	A	118	-43.779	101.670	-15.503	1.00	28.55	T8
	ATOM	8939	CB	LEU	A	118	-43.583	101.466	-17.001	1.00	34.92	T8
	ATOM	8940	CG	LEU	A	118	-43.522	102.714	-17.870	1.00	32.22	T8
	ATOM	8941	CD1	LEU	A	118	-43.399	102.307	-19.321	1.00	30.49	T8
50	ATOM	8942	CD2	LEU	A	118	-44.765	103.543	-17.657	1.00	31.23	T8
	ATOM	8943	C	LEU	A	118	-43.921	100.318	-14.842	1.00	32.92	T8
	ATOM	8944	O	LEU	A	118	-42.926	99.691	-14.490	1.00	25.14	T8
	ATOM	8945	N	GLN	A	119	-45.159	99.873	-14.663	1.00	33.63	T8
	ATOM	8946	CA	GLN	A	119	-45.407	98.573	-14.056	1.00	26.21	T8
55	ATOM	8947	CB	GLN	A	119	-45.453	98.704	-12.542	1.00	36.84	T8
	ATOM	8948	CG	GLN	A	119	-46.571	99.578	-12.045	1.00	27.26	T8
	ATOM	8949	CD	GLN	A	119	-46.529	99.768	-10.541	1.00	27.25	T8
	ATOM	8950	OE1	GLN	A	119	-47.520	100.170	-9.927	1.00	34.16	T8
	ATOM	8951	NE2	GLN	A	119	-45.376	99.491	-9.940	1.00	31.70	T8
60	ATOM	8952	C	GLN	A	119	-46.704	97.959	-14.578	1.00	29.81	T8
	ATOM	8953	O	GLN	A	119	-47.597	98.673	-15.033	1.00	31.59	T8
	ATOM	8954	N	LEU	A	120	-46.781	96.630	-14.524	1.00	27.23	T8
	ATOM	8955	CA	LEU	A	120	-47.939	95.864	-14.987	1.00	30.76	T8
	ATOM	8956	CB	LEU	A	120	-47.452	94.740	-15.902	1.00	35.65	T8
65	ATOM	8957	CG	LEU	A	120	-48.359	93.884	-16.793	1.00	36.15	T8
	ATOM	8958	CD1	LEU	A	120	-49.578	93.418	-16.042	1.00	25.03	T8

	ATOM	8959	CD2	LEU	A	120	-48.760	94.687	-17.988	1.00	25.53	T8
	ATOM	8960	C	LEU	A	120	-48.603	95.274	-13.745	1.00	31.82	T8
	ATOM	8961	O	LEU	A	120	-47.972	94.521	-13.005	1.00	33.79	T8
	ATOM	8962	N	ALA	A	121	-49.871	95.597	-13.514	1.00	25.21	T8
5	ATOM	8963	CA	ALA	A	121	-50.547	95.094	-12.325	1.00	29.72	T8
	ATOM	8964	CB	ALA	A	121	-50.726	96.228	-11.330	1.00	30.37	T8
	ATOM	8965	C	ALA	A	121	-51.886	94.413	-12.564	1.00	29.94	T8
	ATOM	8966	O	ALA	A	121	-52.676	94.845	-13.391	1.00	29.99	T8
	ATOM	8967	N	ILE	A	122	-52.131	93.338	-11.823	1.00	32.38	T8
10	ATOM	8968	CA	ILE	A	122	-53.384	92.602	-11.916	1.00	31.13	T8
	ATOM	8969	CB	ILE	A	122	-53.147	91.100	-12.036	1.00	30.68	T8
	ATOM	8970	CG2	ILE	A	122	-54.476	90.390	-12.167	1.00	31.32	T8
	ATOM	8971	CG1	ILE	A	122	-52.273	90.812	-13.256	1.00	33.56	T8
	ATOM	8972	CD1	ILE	A	122	-51.973	89.357	-13.467	1.00	34.85	T8
15	ATOM	8973	C	ILE	A	122	-54.189	92.878	-10.650	1.00	37.08	T8
	ATOM	8974	O	ILE	A	122	-53.756	92.563	-9.543	1.00	34.68	T8
	ATOM	8975	N	PRO	A	123	-55.372	93.484	-10.801	1.00	29.51	T8
	ATOM	8976	CD	PRO	A	123	-55.888	94.007	-12.071	1.00	32.33	T8
	ATOM	8977	CA	PRO	A	123	-56.276	93.832	-9.700	1.00	35.78	T8
20	ATOM	8978	CB	PRO	A	123	-57.294	94.763	-10.359	1.00	37.49	T8
	ATOM	8979	CG	PRO	A	123	-56.604	95.234	-11.611	1.00	29.46	T8
	ATOM	8980	C	PRO	A	123	-56.966	92.634	-9.056	1.00	29.78	T8
	ATOM	8981	O	PRO	A	123	-58.190	92.591	-8.984	1.00	31.43	T8
	ATOM	8982	N	ARG	A	124	-56.187	91.660	-8.601	1.00	28.72	T8
25	ATOM	8983	CA	ARG	A	124	-56.735	90.477	-7.949	1.00	37.23	T8
	ATOM	8984	CB	ARG	A	124	-56.998	89.372	-8.943	1.00	33.05	T8
	ATOM	8985	CG	ARG	A	124	-58.344	89.475	-9.592	1.00	32.31	T8
	ATOM	8986	CD	ARG	A	124	-58.968	88.099	-9.641	1.00	28.80	T8
	ATOM	8987	NE	ARG	A	124	-59.433	87.650	-8.328	1.00	29.23	T8
30	ATOM	8988	CZ	ARG	A	124	-59.858	86.411	-8.078	1.00	33.38	T8
	ATOM	8989	NH1	ARG	A	124	-59.865	85.505	-9.054	1.00	26.26	T8
	ATOM	8990	NH2	ARG	A	124	-60.288	86.074	-6.863	1.00	26.37	T8
	ATOM	8991	C	ARG	A	124	-55.753	89.990	-6.932	1.00	31.17	T8
	ATOM	8992	O	ARG	A	124	-54.564	90.268	-7.049	1.00	31.52	T8
35	ATOM	8993	N	GLU	A	125	-56.234	89.246	-5.944	1.00	30.54	T8
	ATOM	8994	CA	GLU	A	125	-55.351	88.781	-4.896	1.00	34.56	T8
	ATOM	8995	CB	GLU	A	125	-56.157	88.389	-3.671	1.00	30.01	T8
	ATOM	8996	CG	GLU	A	125	-56.734	89.594	-2.940	1.00	32.52	T8
	ATOM	8997	CD	GLU	A	125	-56.650	89.446	-1.433	1.00	31.77	T8
40	ATOM	8998	OE1	GLU	A	125	-57.179	88.440	-0.897	1.00	31.57	T8
	ATOM	8999	OE2	GLU	A	125	-56.055	90.331	-0.782	1.00	34.75	T8
	ATOM	9000	C	GLU	A	125	-54.392	87.673	-5.298	1.00	35.13	T8
	ATOM	9001	O	GLU	A	125	-53.185	87.774	-5.030	1.00	31.29	T8
	ATOM	9002	N	ASN	A	126	-54.892	86.615	-5.921	1.00	27.82	T8
45	ATOM	9003	CA	ASN	A	126	-53.976	85.573	-6.342	1.00	26.79	T8
	ATOM	9004	CB	ASN	A	126	-53.989	84.386	-5.385	1.00	32.22	T8
	ATOM	9005	CG	ASN	A	126	-53.119	84.615	-4.166	1.00	31.95	T8
	ATOM	9006	OD1	ASN	A	126	-53.519	85.296	-3.223	1.00	32.84	T8
	ATOM	9007	ND2	ASN	A	126	-51.910	84.053	-4.186	1.00	29.46	T8
50	ATOM	9008	C	ASN	A	126	-54.366	85.136	-7.721	1.00	31.36	T8
	ATOM	9009	O	ASN	A	126	-54.753	83.984	-7.943	1.00	28.34	T8
	ATOM	9010	N	ALA	A	127	-54.263	86.081	-8.650	1.00	26.31	T8
	ATOM	9011	CA	ALA	A	127	-54.614	85.846	-10.038	1.00	38.33	T8
	ATOM	9012	CB	ALA	A	127	-54.157	87.020	-10.873	1.00	30.40	T8
55	ATOM	9013	C	ALA	A	127	-54.017	84.551	-10.577	1.00	32.26	T8
	ATOM	9014	O	ALA	A	127	-52.834	84.271	-10.384	1.00	26.83	T8
	ATOM	9015	N	GLN	A	128	-54.854	83.748	-11.225	1.00	35.43	T8
	ATOM	9016	CA	GLN	A	128	-54.392	82.508	-11.827	1.00	26.10	T8
	ATOM	9017	CB	GLN	A	128	-55.548	81.524	-11.946	1.00	29.52	T8
60	ATOM	9018	CG	GLN	A	128	-56.054	81.064	-10.608	1.00	30.07	T8
	ATOM	9019	CD	GLN	A	128	-54.914	80.670	-9.688	1.00	28.47	T8
	ATOM	9020	OE1	GLN	A	128	-54.117	79.788	-10.008	1.00	29.58	T8
	ATOM	9021	NE2	GLN	A	128	-54.825	81.332	-8.540	1.00	27.33	T8
	ATOM	9022	C	GLN	A	128	-53.855	82.902	-13.203	1.00	28.05	T8
65	ATOM	9023	O	GLN	A	128	-54.602	83.091	-14.166	1.00	36.26	T8
	ATOM	9024	N	ILE	A	129	-52.538	83.012	-13.276	1.00	29.19	T8

	ATOM	9025	CA	ILE	A	129	-51.844	83.458	-14.471	1.00	28.38	T8
	ATOM	9026	CB	ILE	A	129	-51.123	84.811	-14.109	1.00	33.56	T8
	ATOM	9027	CG2	ILE	A	129	-49.661	84.803	-14.490	1.00	29.08	T8
	ATOM	9028	CG1	ILE	A	129	-51.882	85.969	-14.727	1.00	29.31	T8
5	ATOM	9029	CD1	ILE	A	129	-53.290	86.061	-14.236	1.00	25.60	T8
	ATOM	9030	C	ILE	A	129	-50.847	82.436	-15.029	1.00	29.02	T8
	ATOM	9031	O	ILE	A	129	-50.414	81.530	-14.323	1.00	39.55	T8
	ATOM	9032	N	SER	A	130	-50.499	82.571	-16.301	1.00	30.94	T8
	ATOM	9033	CA	SER	A	130	-49.511	81.693	-16.907	1.00	35.99	T8
10	ATOM	9034	CB	SER	A	130	-49.927	81.305	-18.317	1.00	24.93	T8
	ATOM	9035	OG	SER	A	130	-48.851	80.684	-18.995	1.00	29.40	T8
	ATOM	9036	C	SER	A	130	-48.212	82.478	-16.970	1.00	30.39	T8
	ATOM	9037	O	SER	A	130	-48.188	83.580	-17.502	1.00	31.09	T8
	ATOM	9038	N	LEU	A	131	-47.131	81.931	-16.435	1.00	31.40	T8
15	ATOM	9039	CA	LEU	A	131	-45.860	82.645	-16.464	1.00	29.03	T8
	ATOM	9040	CB	LEU	A	131	-45.133	82.500	-15.133	1.00	30.20	T8
	ATOM	9041	CG	LEU	A	131	-45.630	83.397	-14.005	1.00	27.12	T8
	ATOM	9042	CD1	LEU	A	131	-47.075	83.126	-13.733	1.00	33.58	T8
	ATOM	9043	CD2	LEU	A	131	-44.823	83.141	-12.770	1.00	34.49	T8
20	ATOM	9044	C	LEU	A	131	-44.922	82.240	-17.590	1.00	34.95	T8
	ATOM	9045	O	LEU	A	131	-43.708	82.270	-17.424	1.00	31.20	T8
	ATOM	9046	N	ASP	A	132	-45.482	81.874	-18.740	1.00	29.19	T8
	ATOM	9047	CA	ASP	A	132	-44.675	81.484	-19.891	1.00	31.80	T8
	ATOM	9048	CB	ASP	A	132	-45.446	80.504	-20.773	1.00	30.83	T8
25	ATOM	9049	CG	ASP	A	132	-45.359	79.075	-20.271	1.00	29.57	T8
	ATOM	9050	OD1	ASP	A	132	-46.122	78.219	-20.773	1.00	34.13	T8
	ATOM	9051	OD2	ASP	A	132	-44.518	78.806	-19.385	1.00	29.13	T8
	ATOM	9052	C	ASP	A	132	-44.234	82.686	-20.719	1.00	30.43	T8
	ATOM	9053	O	ASP	A	132	-44.989	83.644	-20.912	1.00	31.19	T8
30	ATOM	9054	N	GLY	A	133	-42.997	82.616	-21.202	1.00	33.64	T8
	ATOM	9055	CA	GLY	A	133	-42.431	83.684	-22.001	1.00	29.88	T8
	ATOM	9056	C	GLY	A	133	-43.249	84.105	-23.203	1.00	30.46	T8
	ATOM	9057	O	GLY	A	133	-43.202	85.271	-23.577	1.00	34.79	T8
	ATOM	9058	N	ASP	A	134	-43.992	83.182	-23.814	1.00	30.73	T8
35	ATOM	9059	CA	ASP	A	134	-44.800	83.530	-24.979	1.00	28.91	T8
	ATOM	9060	CB	ASP	A	134	-45.419	82.317	-25.656	1.00	35.92	T8
	ATOM	9061	CG	ASP	A	134	-44.667	81.081	-25.430	1.00	34.97	T8
	ATOM	9062	OD1	ASP	A	134	-43.512	81.043	-25.875	1.00	41.42	T8
	ATOM	9063	OD2	ASP	A	134	-45.234	80.152	-24.824	1.00	37.10	T8
40	ATOM	9064	C	ASP	A	134	-45.987	84.366	-24.598	1.00	30.21	T8
	ATOM	9065	O	ASP	A	134	-46.185	85.463	-25.088	1.00	25.98	T8
	ATOM	9066	N	VAL	A	135	-46.801	83.794	-23.732	1.00	31.92	T8
	ATOM	9067	CA	VAL	A	135	-48.034	84.409	-23.314	1.00	25.34	T8
	ATOM	9068	CB	VAL	A	135	-48.885	83.362	-22.619	1.00	32.84	T8
45	ATOM	9069	CG1	VAL	A	135	-49.160	82.233	-23.593	1.00	31.38	T8
	ATOM	9070	CG2	VAL	A	135	-48.159	82.831	-21.392	1.00	37.67	T8
	ATOM	9071	C	VAL	A	135	-48.034	85.697	-22.498	1.00	30.67	T8
	ATOM	9072	O	VAL	A	135	-48.889	86.540	-22.729	1.00	29.67	T8
	ATOM	9073	N	THR	A	136	-47.118	85.875	-21.550	1.00	27.08	T8
50	ATOM	9074	CA	THR	A	136	-47.144	87.129	-20.793	1.00	31.47	T8
	ATOM	9075	CB	THR	A	136	-47.625	86.896	-19.337	1.00	28.85	T8
	ATOM	9076	OG1	THR	A	136	-46.566	86.357	-18.556	1.00	26.15	T8
	ATOM	9077	CG2	THR	A	136	-48.775	85.916	-19.315	1.00	30.22	T8
	ATOM	9078	C	THR	A	136	-45.809	87.904	-20.809	1.00	30.77	T8
55	ATOM	9079	O	THR	A	136	-44.763	87.405	-20.385	1.00	31.73	T8
	ATOM	9080	N	PHE	A	137	-45.866	89.134	-21.315	1.00	23.28	T8
	ATOM	9081	CA	PHE	A	137	-44.690	89.984	-21.439	1.00	28.24	T8
	ATOM	9082	CB	PHE	A	137	-44.055	89.747	-22.806	1.00	33.13	T8
	ATOM	9083	CG	PHE	A	137	-45.041	89.690	-23.936	1.00	32.56	T8
60	ATOM	9084	CD1	PHE	A	137	-45.536	90.848	-24.506	1.00	34.68	T8
	ATOM	9085	CD2	PHE	A	137	-45.462	88.476	-24.437	1.00	40.43	T8
	ATOM	9086	CE1	PHE	A	137	-46.430	90.793	-25.563	1.00	32.12	T8
	ATOM	9087	CE2	PHE	A	137	-46.353	88.414	-25.487	1.00	31.26	T8
	ATOM	9088	CZ	PHE	A	137	-46.837	89.573	-26.049	1.00	26.43	T8
65	ATOM	9089	C	PHE	A	137	-45.010	91.470	-21.238	1.00	28.41	T8
	ATOM	9090	O	PHE	A	137	-46.168	91.871	-21.294	1.00	28.12	T8

	ATOM	9091	N	PHE A 138	-43.980	92.284	-21.018	1.00	30.43	T8
	ATOM	9092	CA	PHE A 138	-44.175	93.704	-20.762	1.00	33.75	T8
	ATOM	9093	CB	PHE A 138	-43.877	93.956	-19.286	1.00	27.80	T8
	ATOM	9094	CG	PHE A 138	-44.321	95.288	-18.782	1.00	27.68	T8
5	ATOM	9095	CD1	PHE A 138	-45.333	95.987	-19.417	1.00	27.63	T8
	ATOM	9096	CD2	PHE A 138	-43.721	95.844	-17.658	1.00	36.77	T8
	ATOM	9097	CE1	PHE A 138	-45.741	97.228	-18.940	1.00	26.95	T8
	ATOM	9098	CE2	PHE A 138	-44.121	97.080	-17.175	1.00	22.16	T8
	ATOM	9099	CZ	PHE A 138	-45.132	97.776	-17.816	1.00	30.18	T8
10	ATOM	9100	C	PHE A 138	-43.325	94.581	-21.686	1.00	28.51	T8
	ATOM	9101	O	PHE A 138	-42.124	94.382	-21.828	1.00	33.31	T8
	ATOM	9102	N	GLY A 139	-43.992	95.571	-22.279	1.00	27.53	T8
	ATOM	9103	CA	GLY A 139	-43.423	96.481	-23.276	1.00	27.26	T8
	ATOM	9104	C	GLY A 139	-42.427	97.598	-23.065	1.00	29.43	T8
15	ATOM	9105	O	GLY A 139	-41.422	97.389	-22.417	1.00	26.10	T8
	ATOM	9106	N	ALA A 140	-42.695	98.757	-23.672	1.00	28.41	T8
	ATOM	9107	CA	ALA A 140	-41.848	99.969	-23.608	1.00	32.03	T8
	ATOM	9108	CB	ALA A 140	-41.224	100.111	-22.234	1.00	35.16	T8
	ATOM	9109	C	ALA A 140	-40.756	100.140	-24.678	1.00	35.64	T8
20	ATOM	9110	O	ALA A 140	-39.706	99.506	-24.631	1.00	34.70	T8
	ATOM	9111	N	LEU A 141	-41.020	101.034	-25.625	1.00	28.56	T8
	ATOM	9112	CA	LEU A 141	-40.105	101.350	-26.726	1.00	31.23	T8
	ATOM	9113	CB	LEU A 141	-40.534	100.615	-28.006	1.00	30.08	T8
	ATOM	9114	CG	LEU A 141	-39.832	100.938	-29.328	1.00	32.60	T8
25	ATOM	9115	CD1	LEU A 141	-40.103	99.856	-30.328	1.00	36.09	T8
	ATOM	9116	CD2	LEU A 141	-40.326	102.251	-29.867	1.00	32.39	T8
	ATOM	9117	C	LEU A 141	-40.148	102.865	-26.955	1.00	31.04	T8
	ATOM	9118	O	LEU A 141	-41.223	103.463	-26.999	1.00	36.45	T8
	ATOM	9119	N	LYS A 142	-38.988	103.487	-27.120	1.00	31.06	T8
30	ATOM	9120	CA	LYS A 142	-38.946	104.930	-27.311	1.00	35.62	T8
	ATOM	9121	CB	LYS A 142	-37.657	105.501	-26.720	1.00	28.52	T8
	ATOM	9122	CG	LYS A 142	-37.572	107.005	-26.863	1.00	40.77	T8
	ATOM	9123	CD	LYS A 142	-36.466	107.618	-26.036	1.00	31.00	T8
	ATOM	9124	CE	LYS A 142	-36.460	109.125	-26.231	1.00	36.55	T8
35	ATOM	9125	NZ	LYS A 142	-35.369	109.776	-25.470	1.00	34.37	T8
	ATOM	9126	C	LYS A 142	-39.092	105.415	-28.747	1.00	26.14	T8
	ATOM	9127	O	LYS A 142	-38.374	104.970	-29.637	1.00	34.92	T8
	ATOM	9128	N	LEU A 143	-40.015	106.348	-28.963	1.00	32.72	T8
	ATOM	9129	CA	LEU A 143	-40.246	106.907	-30.291	1.00	31.07	T8
40	ATOM	9130	CB	LEU A 143	-41.652	107.504	-30.376	1.00	24.50	T8
	ATOM	9131	CG	LEU A 143	-42.852	106.596	-30.095	1.00	27.20	T8
	ATOM	9132	CD1	LEU A 143	-44.125	107.426	-30.147	1.00	28.49	T8
	ATOM	9133	CD2	LEU A 143	-42.914	105.471	-31.111	1.00	36.64	T8
	ATOM	9134	C	LEU A 143	-39.229	108.003	-30.593	1.00	25.83	T8
45	ATOM	9135	O	LEU A 143	-38.682	108.622	-29.674	1.00	29.37	T8
	ATOM	9136	N	LEU A 144	-38.979	108.247	-31.878	1.00	27.02	T8
	ATOM	9137	CA	LEU A 144	-38.034	109.285	-32.268	1.00	33.05	T8
	ATOM	9138	CB	LEU A 144	-37.393	108.954	-33.609	1.00	33.90	T8
	ATOM	9139	CG	LEU A 144	-36.491	107.721	-33.634	1.00	27.68	T8
50	ATOM	9140	CD1	LEU A 144	-36.092	107.422	-35.065	1.00	22.18	T8
	ATOM	9141	CD2	LEU A 144	-35.255	107.944	-32.773	1.00	35.14	T8
	ATOM	9142	C	LEU A 144	-38.742	110.619	-32.369	1.00	34.04	T8
	ATOM	9143	O	LEU A 144	-39.989	110.625	-32.453	1.00	34.01	T8
	ATOM	9144	OXT	LEU A 144	-38.035	111.645	-32.376	1.00	28.77	T8
55	ATOM	9145	CB	VAL A 1	-36.000	115.069	-37.089	1.00	39.15	T9
	ATOM	9146	CG1	VAL A 1	-35.250	115.193	-35.754	1.00	26.43	T9
	ATOM	9147	CG2	VAL A 1	-35.225	115.770	-38.195	1.00	27.00	T9
	ATOM	9148	C	VAL A 1	-37.108	112.910	-36.385	1.00	31.42	T9
	ATOM	9149	O	VAL A 1	-36.665	111.984	-35.705	1.00	29.56	T9
60	ATOM	9150	N	VAL A 1	-36.778	113.411	-38.830	1.00	28.58	T9
	ATOM	9151	CA	VAL A 1	-36.212	113.561	-37.455	1.00	41.46	T9
	ATOM	9152	N	THR A 2	-38.351	113.382	-36.235	1.00	34.96	T9
	ATOM	9153	CA	THR A 2	-39.251	112.814	-35.227	1.00	26.32	T9
	ATOM	9154	CB	THR A 2	-39.809	113.895	-34.272	1.00	24.19	T9
65	ATOM	9155	OG1	THR A 2	-40.754	114.707	-34.977	1.00	32.27	T9
	ATOM	9156	CG2	THR A 2	-38.685	114.760	-33.725	1.00	39.07	T9

	ATOM	9157	C	THR	A	2	-40.440	112.080	-35.829	1.00	32.66	T9
	ATOM	9158	O	THR	A	2	-40.664	112.142	-37.032	1.00	34.78	T9
	ATOM	9159	N	GLN	A	3	-41.207	111.396	-34.980	1.00	29.52	T9
	ATOM	9160	CA	GLN	A	3	-42.382	110.652	-35.425	1.00	31.52	T9
5	ATOM	9161	CB	GLN	A	3	-42.300	109.209	-34.942	1.00	30.50	T9
	ATOM	9162	CG	GLN	A	3	-40.956	108.559	-35.153	1.00	33.12	T9
	ATOM	9163	CD	GLN	A	3	-40.980	107.092	-34.793	1.00	20.25	T9
	ATOM	9164	OE1	GLN	A	3	-41.720	106.321	-35.395	1.00	28.02	T9
	ATOM	9165	NE2	GLN	A	3	-40.176	106.696	-33.804	1.00	27.07	T9
10	ATOM	9166	C	GLN	A	3	-43.682	111.266	-34.905	1.00	31.62	T9
	ATOM	9167	O	GLN	A	3	-44.025	111.107	-33.732	1.00	29.25	T9
	ATOM	9168	N	ASP	A	4	-44.411	111.961	-35.767	1.00	24.46	T9
	ATOM	9169	CA	ASP	A	4	-45.663	112.559	-35.338	1.00	31.05	T9
	ATOM	9170	CB	ASP	A	4	-46.299	113.346	-36.478	1.00	35.15	T9
15	ATOM	9171	CG	ASP	A	4	-45.450	114.518	-36.916	1.00	26.03	T9
	ATOM	9172	OD1	ASP	A	4	-44.550	114.904	-36.138	1.00	30.43	T9
	ATOM	9173	OD2	ASP	A	4	-45.685	115.061	-38.023	1.00	25.30	T9
	ATOM	9174	C	ASP	A	4	-46.619	111.476	-34.881	1.00	33.59	T9
	ATOM	9175	O	ASP	A	4	-46.588	110.357	-35.390	1.00	35.72	T9
20	ATOM	9176	N	CYS	A	5	-47.460	111.813	-33.911	1.00	31.69	T9
	ATOM	9177	CA	CYS	A	5	-48.450	110.883	-33.378	1.00	36.37	T9
	ATOM	9178	CB	CYS	A	5	-47.813	109.917	-32.358	1.00	42.70	T9
	ATOM	9179	SG	CYS	A	5	-46.391	110.544	-31.380	1.00	34.30	T9
	ATOM	9180	C	CYS	A	5	-49.588	111.655	-32.730	1.00	28.86	T9
25	ATOM	9181	O	CYS	A	5	-49.385	112.737	-32.193	1.00	33.15	T9
	ATOM	9182	N	LEU	A	6	-50.797	111.115	-32.812	1.00	34.15	T9
	ATOM	9183	CA	LEU	A	6	-51.962	111.752	-32.210	1.00	24.09	T9
	ATOM	9184	CB	LEU	A	6	-52.784	112.478	-33.273	1.00	31.31	T9
	ATOM	9185	CG	LEU	A	6	-54.071	113.154	-32.799	1.00	26.35	T9
30	ATOM	9186	CD1	LEU	A	6	-54.375	114.360	-33.657	1.00	35.02	T9
	ATOM	9187	CD2	LEU	A	6	-55.213	112.166	-32.844	1.00	35.06	T9
	ATOM	9188	C	LEU	A	6	-52.781	110.659	-31.554	1.00	30.02	T9
	ATOM	9189	O	LEU	A	6	-52.999	109.614	-32.150	1.00	33.02	T9
	ATOM	9190	N	GLN	A	7	-53.224	110.888	-30.324	1.00	35.96	T9
35	ATOM	9191	CA	GLN	A	7	-53.996	109.881	-29.618	1.00	29.61	T9
	ATOM	9192	CB	GLN	A	7	-53.147	109.272	-28.512	1.00	29.45	T9
	ATOM	9193	CG	GLN	A	7	-53.742	108.042	-27.872	1.00	29.90	T9
	ATOM	9194	CD	GLN	A	7	-52.755	107.365	-26.955	1.00	25.26	T9
	ATOM	9195	OE1	GLN	A	7	-52.418	107.886	-25.897	1.00	30.31	T9
40	ATOM	9196	NE2	GLN	A	7	-52.264	106.206	-27.367	1.00	24.95	T9
	ATOM	9197	C	GLN	A	7	-55.277	110.442	-29.035	1.00	26.24	T9
	ATOM	9198	O	GLN	A	7	-55.297	111.556	-28.526	1.00	25.51	T9
	ATOM	9199	N	LEU	A	8	-56.345	109.657	-29.110	1.00	33.12	T9
	ATOM	9200	CA	LEU	A	8	-57.640	110.062	-28.592	1.00	26.81	T9
45	ATOM	9201	CB	LEU	A	8	-58.680	110.015	-29.708	1.00	25.76	T9
	ATOM	9202	CG	LEU	A	8	-58.881	111.212	-30.636	1.00	30.34	T9
	ATOM	9203	CD1	LEU	A	8	-57.739	112.186	-30.523	1.00	26.32	T9
	ATOM	9204	CD2	LEU	A	8	-59.038	110.709	-32.042	1.00	28.08	T9
	ATOM	9205	C	LEU	A	8	-58.089	109.180	-27.435	1.00	36.86	T9
50	ATOM	9206	O	LEU	A	8	-57.692	108.027	-27.325	1.00	30.53	T9
	ATOM	9207	N	ILE	A	9	-58.928	109.743	-26.577	1.00	33.62	T9
	ATOM	9208	CA	ILE	A	9	-59.452	109.053	-25.402	1.00	22.76	T9
	ATOM	9209	CB	ILE	A	9	-58.898	109.684	-24.109	1.00	28.34	T9
	ATOM	9210	CG2	ILE	A	9	-59.759	109.327	-22.930	1.00	32.67	T9
55	ATOM	9211	CG1	ILE	A	9	-57.483	109.217	-23.850	1.00	31.49	T9
	ATOM	9212	CD1	ILE	A	9	-56.918	109.869	-22.617	1.00	24.58	T9
	ATOM	9213	C	ILE	A	9	-60.971	109.206	-25.362	1.00	31.21	T9
	ATOM	9214	O	ILE	A	9	-61.498	110.250	-25.736	1.00	27.59	T9
	ATOM	9215	N	ALA	A	10	-61.673	108.185	-24.890	1.00	32.82	T9
60	ATOM	9216	CA	ALA	A	10	-63.120	108.275	-24.808	1.00	23.79	T9
	ATOM	9217	CB	ALA	A	10	-63.698	106.969	-24.323	1.00	32.71	T9
	ATOM	9218	C	ALA	A	10	-63.542	109.399	-23.870	1.00	32.41	T9
	ATOM	9219	O	ALA	A	10	-63.044	109.508	-22.748	1.00	33.50	T9
	ATOM	9220	N	ASP	A	11	-64.462	110.238	-24.334	1.00	25.73	T9
65	ATOM	9221	CA	ASP	A	11	-64.963	111.340	-23.522	1.00	35.30	T9
	ATOM	9222	CB	ASP	A	11	-65.306	112.541	-24.395	1.00	30.60	T9

	ATOM	9223	CG	ASP	A	11	-65.968	113.644	-23.609	1.00	26.84	T9
	ATOM	9224	OD1	ASP	A	11	-65.545	113.864	-22.457	1.00	31.40	T9
	ATOM	9225	OD2	ASP	A	11	-66.900	114.290	-24.131	1.00	30.81	T9
	ATOM	9226	C	ASP	A	11	-66.197	110.890	-22.754	1.00	25.29	T9
5	ATOM	9227	O	ASP	A	11	-67.327	110.996	-23.230	1.00	32.78	T9
	ATOM	9228	N	SER	A	12	-65.947	110.380	-21.554	1.00	31.19	T9
	ATOM	9229	CA	SER	A	12	-66.983	109.869	-20.657	1.00	26.56	T9
	ATOM	9230	CB	SER	A	12	-66.338	109.232	-19.413	1.00	32.31	T9
	ATOM	9231	OG	SER	A	12	-65.520	110.153	-18.697	1.00	27.92	T9
10	ATOM	9232	C	SER	A	12	-67.978	110.924	-20.208	1.00	33.86	T9
	ATOM	9233	O	SER	A	12	-68.620	110.767	-19.179	1.00	33.32	T9
	ATOM	9234	N	GLU	A	13	-68.111	112.001	-20.968	1.00	28.94	T9
	ATOM	9235	CA	GLU	A	13	-69.048	113.037	-20.579	1.00	33.23	T9
	ATOM	9236	CB	GLU	A	13	-68.315	114.163	-19.871	1.00	24.70	T9
15	ATOM	9237	CG	GLU	A	13	-68.241	113.924	-18.392	1.00	32.20	T9
	ATOM	9238	CD	GLU	A	13	-67.605	115.079	-17.676	1.00	33.59	T9
	ATOM	9239	OE1	GLU	A	13	-67.937	116.240	-18.038	1.00	32.79	T9
	ATOM	9240	OE2	GLU	A	13	-66.779	114.828	-16.757	1.00	28.03	T9
	ATOM	9241	C	GLU	A	13	-69.926	113.594	-21.682	1.00	28.89	T9
20	ATOM	9242	O	GLU	A	13	-70.409	114.723	-21.599	1.00	24.88	T9
	ATOM	9243	N	THR	A	14	-70.120	112.801	-22.722	1.00	32.11	T9
	ATOM	9244	CA	THR	A	14	-70.995	113.185	-23.805	1.00	31.49	T9
	ATOM	9245	CB	THR	A	14	-70.262	113.900	-24.947	1.00	33.22	T9
	ATOM	9246	OG1	THR	A	14	-69.222	113.061	-25.431	1.00	25.89	T9
25	ATOM	9247	CG2	THR	A	14	-69.676	115.216	-24.472	1.00	32.27	T9
	ATOM	9248	C	THR	A	14	-71.581	111.882	-24.304	1.00	36.99	T9
	ATOM	9249	O	THR	A	14	-70.961	110.826	-24.188	1.00	29.92	T9
	ATOM	9250	N	PRO	A	15	-72.799	111.936	-24.845	1.00	28.77	T9
	ATOM	9251	CD	PRO	A	15	-73.608	113.145	-25.054	1.00	36.30	T9
30	ATOM	9252	CA	PRO	A	15	-73.487	110.753	-25.362	1.00	31.85	T9
	ATOM	9253	CB	PRO	A	15	-74.835	111.310	-25.828	1.00	28.70	T9
	ATOM	9254	CG	PRO	A	15	-74.993	112.580	-25.031	1.00	28.60	T9
	ATOM	9255	C	PRO	A	15	-72.718	110.124	-26.511	1.00	31.30	T9
	ATOM	9256	O	PRO	A	15	-72.082	110.825	-27.289	1.00	32.86	T9
35	ATOM	9257	N	THR	A	16	-72.783	108.808	-26.616	1.00	32.10	T9
	ATOM	9258	CA	THR	A	16	-72.108	108.127	-27.702	1.00	34.59	T9
	ATOM	9259	CB	THR	A	16	-72.097	106.629	-27.454	1.00	31.61	T9
	ATOM	9260	OG1	THR	A	16	-73.428	106.110	-27.586	1.00	29.92	T9
	ATOM	9261	CG2	THR	A	16	-71.609	106.351	-26.053	1.00	25.33	T9
40	ATOM	9262	C	THR	A	16	-72.870	108.416	-28.998	1.00	27.80	T9
	ATOM	9263	O	THR	A	16	-74.027	108.034	-29.138	1.00	37.38	T9
	ATOM	9264	N	ILE	A	17	-72.220	109.085	-29.942	1.00	31.91	T9
	ATOM	9265	CA	ILE	A	17	-72.852	109.440	-31.210	1.00	24.55	T9
	ATOM	9266	CB	ILE	A	17	-71.836	110.091	-32.153	1.00	32.34	T9
45	ATOM	9267	CG2	ILE	A	17	-72.495	110.429	-33.469	1.00	35.08	T9
	ATOM	9268	CG1	ILE	A	17	-71.269	111.350	-31.501	1.00	34.60	T9
	ATOM	9269	CD1	ILE	A	17	-70.200	112.031	-32.311	1.00	29.01	T9
	ATOM	9270	C	ILE	A	17	-73.525	108.284	-31.953	1.00	27.31	T9
	ATOM	9271	O	ILE	A	17	-72.930	107.221	-32.139	1.00	24.40	T9
50	ATOM	9272	N	GLN	A	18	-74.770	108.508	-32.378	1.00	31.70	T9
	ATOM	9273	CA	GLN	A	18	-75.538	107.506	-33.120	1.00	28.18	T9
	ATOM	9274	CB	GLN	A	18	-76.871	107.239	-32.446	1.00	37.62	T9
	ATOM	9275	CG	GLN	A	18	-77.067	105.790	-32.127	1.00	28.04	T9
	ATOM	9276	CD	GLN	A	18	-76.212	105.355	-30.967	1.00	25.94	T9
55	ATOM	9277	OE1	GLN	A	18	-76.526	105.641	-29.810	1.00	30.30	T9
	ATOM	9278	NE2	GLN	A	18	-75.109	104.669	-31.264	1.00	33.75	T9
	ATOM	9279	C	GLN	A	18	-75.793	107.964	-34.553	1.00	26.72	T9
	ATOM	9280	O	GLN	A	18	-76.183	109.107	-34.789	1.00	34.50	T9
	ATOM	9281	N	LYS	A	19	-75.585	107.071	-35.513	1.00	35.59	T9
60	ATOM	9282	CA	LYS	A	19	-75.778	107.426	-36.914	1.00	26.47	T9
	ATOM	9283	CB	LYS	A	19	-74.753	108.477	-37.321	1.00	32.68	T9
	ATOM	9284	CG	LYS	A	19	-74.694	108.727	-38.809	1.00	40.59	T9
	ATOM	9285	CD	LYS	A	19	-73.821	109.941	-39.119	1.00	28.23	T9
	ATOM	9286	CE	LYS	A	19	-73.745	110.205	-40.625	1.00	28.59	T9
65	ATOM	9287	NZ	LYS	A	19	-72.939	111.426	-40.961	1.00	33.34	T9
	ATOM	9288	C	LYS	A	19	-75.672	106.226	-37.848	1.00	32.73	T9

	ATOM	9289	O	LYS	A	19	-74.710	105.458	-37.778	1.00	30.36	T9
	ATOM	9290	N	GLY	A	20	-76.658	106.084	-38.732	1.00	28.60	T9
	ATOM	9291	CA	GLY	A	20	-76.677	104.969	-39.667	1.00	30.84	T9
	ATOM	9292	C	GLY	A	20	-76.576	103.652	-38.923	1.00	37.45	T9
5	ATOM	9293	O	GLY	A	20	-75.868	102.744	-39.362	1.00	28.78	T9
	ATOM	9294	N	SER	A	21	-77.281	103.561	-37.792	1.00	26.59	T9
	ATOM	9295	CA	SER	A	21	-77.279	102.360	-36.945	1.00	30.54	T9
	ATOM	9296	CB	SER	A	21	-78.071	101.224	-37.610	1.00	27.73	T9
	ATOM	9297	OG	SER	A	21	-77.507	100.859	-38.860	1.00	29.23	T9
10	ATOM	9298	C	SER	A	21	-75.857	101.886	-36.597	1.00	30.84	T9
	ATOM	9299	O	SER	A	21	-75.560	100.685	-36.563	1.00	34.02	T9
	ATOM	9300	N	TYR	A	22	-74.993	102.866	-36.341	1.00	31.60	T9
	ATOM	9301	CA	TYR	A	22	-73.603	102.648	-35.961	1.00	34.96	T9
	ATOM	9302	CB	TYR	A	22	-72.667	103.146	-37.061	1.00	28.14	T9
15	ATOM	9303	CG	TYR	A	22	-72.216	102.088	-38.030	1.00	33.17	T9
	ATOM	9304	CD1	TYR	A	22	-72.979	100.944	-38.251	1.00	28.62	T9
	ATOM	9305	CE1	TYR	A	22	-72.573	99.972	-39.170	1.00	29.51	T9
	ATOM	9306	CD2	TYR	A	22	-71.036	102.240	-38.752	1.00	29.62	T9
	ATOM	9307	CE2	TYR	A	22	-70.622	101.278	-39.677	1.00	28.93	T9
20	ATOM	9308	CZ	TYR	A	22	-71.394	100.149	-39.877	1.00	25.80	T9
	ATOM	9309	OH	TYR	A	22	-70.984	99.195	-40.777	1.00	32.83	T9
	ATOM	9310	C	TYR	A	22	-73.373	103.472	-34.702	1.00	30.83	T9
	ATOM	9311	O	TYR	A	22	-73.977	104.531	-34.537	1.00	33.06	T9
	ATOM	9312	N	THR	A	23	-72.519	102.992	-33.808	1.00	27.28	T9
25	ATOM	9313	CA	THR	A	23	-72.233	103.751	-32.607	1.00	26.98	T9
	ATOM	9314	CB	THR	A	23	-72.323	102.890	-31.341	1.00	31.91	T9
	ATOM	9315	OG1	THR	A	23	-73.450	102.011	-31.423	1.00	27.52	T9
	ATOM	9316	CG2	THR	A	23	-72.505	103.779	-30.135	1.00	34.92	T9
	ATOM	9317	C	THR	A	23	-70.818	104.303	-32.727	1.00	31.92	T9
30	ATOM	9318	O	THR	A	23	-69.883	103.574	-33.061	1.00	25.76	T9
	ATOM	9319	N	PHE	A	24	-70.673	105.598	-32.470	1.00	25.02	T9
	ATOM	9320	CA	PHE	A	24	-69.374	106.249	-32.543	1.00	28.55	T9
	ATOM	9321	CB	PHE	A	24	-69.402	107.364	-33.583	1.00	28.74	T9
	ATOM	9322	CG	PHE	A	24	-69.593	106.874	-34.984	1.00	28.77	T9
35	ATOM	9323	CD1	PHE	A	24	-70.853	106.551	-35.457	1.00	31.78	T9
	ATOM	9324	CD2	PHE	A	24	-68.501	106.717	-35.828	1.00	40.33	T9
	ATOM	9325	CE1	PHE	A	24	-71.023	106.078	-36.750	1.00	30.09	T9
	ATOM	9326	CE2	PHE	A	24	-68.663	106.244	-37.118	1.00	32.57	T9
	ATOM	9327	CZ	PHE	A	24	-69.929	105.925	-37.579	1.00	31.50	T9
40	ATOM	9328	C	PHE	A	24	-68.951	106.816	-31.193	1.00	34.74	T9
	ATOM	9329	O	PHE	A	24	-69.674	107.597	-30.576	1.00	31.85	T9
	ATOM	9330	N	VAL	A	25	-67.770	106.408	-30.741	1.00	33.13	T9
	ATOM	9331	CA	VAL	A	25	-67.223	106.859	-29.470	1.00	26.31	T9
	ATOM	9332	CB	VAL	A	25	-65.924	106.123	-29.140	1.00	25.64	T9
45	ATOM	9333	CG1	VAL	A	25	-65.337	106.657	-27.853	1.00	30.64	T9
	ATOM	9334	CG2	VAL	A	25	-66.188	104.643	-29.029	1.00	30.17	T9
	ATOM	9335	C	VAL	A	25	-66.914	108.344	-29.507	1.00	31.29	T9
	ATOM	9336	O	VAL	A	25	-66.325	108.840	-30.459	1.00	34.50	T9
	ATOM	9337	N	PRO	A	26	-67.325	109.081	-28.469	1.00	27.36	T9
50	ATOM	9338	CD	PRO	A	26	-68.211	108.635	-27.383	1.00	29.10	T9
	ATOM	9339	CA	PRO	A	26	-67.081	110.525	-28.385	1.00	31.60	T9
	ATOM	9340	CB	PRO	A	26	-68.018	110.972	-27.270	1.00	30.36	T9
	ATOM	9341	CG	PRO	A	26	-69.030	109.856	-27.158	1.00	24.91	T9
	ATOM	9342	C	PRO	A	26	-65.622	110.703	-27.984	1.00	32.72	T9
55	ATOM	9343	O	PRO	A	26	-65.245	110.376	-26.868	1.00	37.21	T9
	ATOM	9344	N	TRP	A	27	-64.795	111.214	-28.882	1.00	31.29	T9
	ATOM	9345	CA	TRP	A	27	-63.386	111.377	-28.558	1.00	34.50	T9
	ATOM	9346	CB	TRP	A	27	-62.530	111.243	-29.821	1.00	36.74	T9
	ATOM	9347	CG	TRP	A	27	-62.619	109.902	-30.439	1.00	24.60	T9
60	ATOM	9348	CD2	TRP	A	27	-62.380	108.642	-29.800	1.00	23.84	T9
	ATOM	9349	CE2	TRP	A	27	-62.643	107.636	-30.749	1.00	29.67	T9
	ATOM	9350	CE3	TRP	A	27	-61.975	108.265	-28.517	1.00	31.68	T9
	ATOM	9351	CD1	TRP	A	27	-62.997	109.619	-31.710	1.00	34.42	T9
	ATOM	9352	NE1	TRP	A	27	-63.018	108.259	-31.908	1.00	27.60	T9
65	ATOM	9353	CZ2	TRP	A	27	-62.514	106.278	-30.457	1.00	37.11	T9
	ATOM	9354	CZ3	TRP	A	27	-61.849	106.912	-28.228	1.00	33.95	T9

	ATOM	9355	CH2	TRP	A	27	-62.119	105.939	-29.195	1.00	29.14	T9
	ATOM	9356	C	TRP	A	27	-62.999	112.659	-27.829	1.00	32.68	T9
	ATOM	9357	O	TRP	A	27	-63.748	113.639	-27.788	1.00	26.82	T9
	ATOM	9358	N	LEU	A	28	-61.805	112.619	-27.247	1.00	30.72	T9
5	ATOM	9359	CA	LEU	A	28	-61.241	113.734	-26.516	1.00	31.86	T9
	ATOM	9360	CB	LEU	A	28	-61.613	113.623	-25.052	1.00	30.87	T9
	ATOM	9361	CG	LEU	A	28	-61.494	114.925	-24.270	1.00	35.21	T9
	ATOM	9362	CD1	LEU	A	28	-62.480	115.943	-24.846	1.00	26.80	T9
	ATOM	9363	CD2	LEU	A	28	-61.779	114.652	-22.794	1.00	34.60	T9
10	ATOM	9364	C	LEU	A	28	-59.734	113.616	-26.691	1.00	30.24	T9
	ATOM	9365	O	LEU	A	28	-59.169	112.548	-26.474	1.00	29.20	T9
	ATOM	9366	N	LEU	A	29	-59.083	114.701	-27.092	1.00	28.04	T9
	ATOM	9367	CA	LEU	A	29	-57.650	114.655	-27.322	1.00	28.77	T9
	ATOM	9368	CB	LEU	A	29	-57.119	116.026	-27.723	1.00	25.40	T9
15	ATOM	9369	CG	LEU	A	29	-55.608	116.027	-27.942	1.00	27.94	T9
	ATOM	9370	CD1	LEU	A	29	-55.291	115.376	-29.264	1.00	31.76	T9
	ATOM	9371	CD2	LEU	A	29	-55.088	117.435	-27.914	1.00	28.95	T9
	ATOM	9372	C	LEU	A	29	-56.875	114.178	-26.123	1.00	29.00	T9
	ATOM	9373	O	LEU	A	29	-57.021	114.721	-25.033	1.00	28.76	T9
20	ATOM	9374	N	SER	A	30	-56.057	113.153	-26.328	1.00	36.03	T9
	ATOM	9375	CA	SER	A	30	-55.216	112.633	-25.267	1.00	30.32	T9
	ATOM	9376	CB	SER	A	30	-54.892	111.164	-25.499	1.00	37.68	T9
	ATOM	9377	OG	SER	A	30	-53.976	110.706	-24.532	1.00	25.96	T9
	ATOM	9378	C	SER	A	30	-53.949	113.459	-25.372	1.00	29.18	T9
25	ATOM	9379	O	SER	A	30	-53.520	114.087	-24.416	1.00	26.62	T9
	ATOM	9380	N	PHE	A	31	-53.362	113.467	-26.558	1.00	32.93	T9
	ATOM	9381	CA	PHE	A	31	-52.155	114.235	-26.808	1.00	31.81	T9
	ATOM	9382	CB	PHE	A	31	-50.955	113.611	-26.086	1.00	43.62	T9
	ATOM	9383	CG	PHE	A	31	-50.262	112.535	-26.866	1.00	30.58	T9
30	ATOM	9384	CD1	PHE	A	31	-49.306	112.857	-27.823	1.00	22.83	T9
	ATOM	9385	CD2	PHE	A	31	-50.568	111.194	-26.652	1.00	34.22	T9
	ATOM	9386	CE1	PHE	A	31	-48.664	111.861	-28.554	1.00	34.98	T9
	ATOM	9387	CE2	PHE	A	31	-49.933	110.193	-27.378	1.00	42.04	T9
	ATOM	9388	CZ	PHE	A	31	-48.978	110.528	-28.331	1.00	32.25	T9
35	ATOM	9389	C	PHE	A	31	-51.914	114.270	-28.310	1.00	35.43	T9
	ATOM	9390	O	PHE	A	31	-52.347	113.382	-29.041	1.00	32.60	T9
	ATOM	9391	N	LYS	A	32	-51.235	115.314	-28.766	1.00	31.66	T9
	ATOM	9392	CA	LYS	A	32	-50.931	115.476	-30.178	1.00	32.41	T9
	ATOM	9393	CB	LYS	A	32	-51.845	116.530	-30.795	1.00	34.27	T9
40	ATOM	9394	CG	LYS	A	32	-51.396	116.945	-32.161	1.00	21.55	T9
	ATOM	9395	CD	LYS	A	32	-52.149	118.140	-32.680	1.00	33.18	T9
	ATOM	9396	CE	LYS	A	32	-51.456	118.652	-33.929	1.00	30.02	T9
	ATOM	9397	NZ	LYS	A	32	-52.252	119.715	-34.581	1.00	35.94	T9
	ATOM	9398	C	LYS	A	32	-49.477	115.906	-30.310	1.00	31.58	T9
45	ATOM	9399	O	LYS	A	32	-49.048	116.880	-29.700	1.00	32.67	T9
	ATOM	9400	N	ARG	A	33	-48.720	115.176	-31.113	1.00	31.22	T9
	ATOM	9401	CA	ARG	A	33	-47.309	115.466	-31.294	1.00	27.38	T9
	ATOM	9402	CB	ARG	A	33	-46.484	114.358	-30.632	1.00	26.96	T9
	ATOM	9403	CG	ARG	A	33	-44.992	114.389	-30.860	1.00	31.67	T9
50	ATOM	9404	CD	ARG	A	33	-44.309	113.585	-29.767	1.00	24.27	T9
	ATOM	9405	NE	ARG	A	33	-42.863	113.477	-29.937	1.00	31.27	T9
	ATOM	9406	CZ	ARG	A	33	-42.279	112.598	-30.741	1.00	30.55	T9
	ATOM	9407	NH1	ARG	A	33	-43.020	111.751	-31.439	1.00	25.45	T9
	ATOM	9408	NH2	ARG	A	33	-40.963	112.575	-30.858	1.00	21.27	T9
55	ATOM	9409	C	ARG	A	33	-46.981	115.561	-32.766	1.00	32.17	T9
	ATOM	9410	O	ARG	A	33	-47.231	114.620	-33.530	1.00	33.16	T9
	ATOM	9411	N	GLY	A	34	-46.429	116.702	-33.167	1.00	38.31	T9
	ATOM	9412	CA	GLY	A	34	-46.070	116.891	-34.558	1.00	25.91	T9
	ATOM	9413	C	GLY	A	34	-47.163	117.559	-35.368	1.00	29.83	T9
60	ATOM	9414	O	GLY	A	34	-48.117	118.123	-34.811	1.00	29.80	T9
	ATOM	9415	N	SER	A	35	-47.040	117.465	-36.690	1.00	34.22	T9
	ATOM	9416	CA	SER	A	35	-47.993	118.094	-37.596	1.00	30.20	T9
	ATOM	9417	CB	SER	A	35	-47.260	119.133	-38.436	1.00	30.56	T9
	ATOM	9418	OG	SER	A	35	-46.158	118.533	-39.108	1.00	22.36	T9
65	ATOM	9419	C	SER	A	35	-48.750	117.162	-38.536	1.00	31.64	T9
	ATOM	9420	O	SER	A	35	-49.802	117.533	-39.047	1.00	33.87	T9

	ATOM	9421	N	ALA	A	36	-48.226	115.963	-38.767	1.00	28.58	T9
	ATOM	9422	CA	ALA	A	36	-48.856	115.026	-39.691	1.00	30.83	T9
	ATOM	9423	CB	ALA	A	36	-47.968	113.812	-39.883	1.00	41.14	T9
	ATOM	9424	C	ALA	A	36	-50.267	114.573	-39.356	1.00	27.35	T9
5	ATOM	9425	O	ALA	A	36	-50.948	114.043	-40.235	1.00	31.30	T9
	ATOM	9426	N	LEU	A	37	-50.715	114.774	-38.114	1.00	30.38	T9
	ATOM	9427	CA	LEU	A	37	-52.058	114.343	-37.720	1.00	36.18	T9
	ATOM	9428	CB	LEU	A	37	-51.964	113.030	-36.936	1.00	24.32	T9
	ATOM	9429	CG	LEU	A	37	-51.351	111.866	-37.720	1.00	26.12	T9
10	ATOM	9430	CD1	LEU	A	37	-50.885	110.782	-36.779	1.00	30.13	T9
	ATOM	9431	CD2	LEU	A	37	-52.365	111.336	-38.705	1.00	30.90	T9
	ATOM	9432	C	LEU	A	37	-52.811	115.384	-36.905	1.00	36.96	T9
	ATOM	9433	O	LEU	A	37	-52.210	116.129	-36.142	1.00	29.69	T9
	ATOM	9434	N	GLU	A	38	-54.132	115.418	-37.067	1.00	28.76	T9
15	ATOM	9435	CA	GLU	A	38	-54.987	116.376	-36.363	1.00	29.72	T9
	ATOM	9436	CB	GLU	A	38	-55.220	117.615	-37.232	1.00	29.51	T9
	ATOM	9437	CG	GLU	A	38	-54.123	118.656	-37.216	1.00	28.40	T9
	ATOM	9438	CD	GLU	A	38	-54.315	119.706	-38.302	1.00	25.77	T9
	ATOM	9439	OE1	GLU	A	38	-55.486	120.063	-38.582	1.00	29.95	T9
20	ATOM	9440	OE2	GLU	A	38	-53.294	120.177	-38.861	1.00	25.94	T9
	ATOM	9441	C	GLU	A	38	-56.359	115.801	-36.021	1.00	30.92	T9
	ATOM	9442	O	GLU	A	38	-56.765	114.785	-36.573	1.00	38.15	T9
	ATOM	9443	N	GLU	A	39	-57.076	116.460	-35.113	1.00	34.55	T9
	ATOM	9444	CA	GLU	A	39	-58.422	116.021	-34.761	1.00	29.93	T9
25	ATOM	9445	CB	GLU	A	39	-58.825	116.421	-33.354	1.00	29.68	T9
	ATOM	9446	CG	GLU	A	39	-57.793	116.239	-32.308	1.00	22.57	T9
	ATOM	9447	CD	GLU	A	39	-57.977	117.257	-31.198	1.00	31.78	T9
	ATOM	9448	OE1	GLU	A	39	-57.268	118.304	-31.235	1.00	24.82	T9
	ATOM	9449	OE2	GLU	A	39	-58.846	117.018	-30.312	1.00	24.09	T9
30	ATOM	9450	C	GLU	A	39	-59.326	116.809	-35.669	1.00	33.47	T9
	ATOM	9451	O	GLU	A	39	-59.040	117.963	-35.989	1.00	31.90	T9
	ATOM	9452	N	LYS	A	40	-60.430	116.205	-36.070	1.00	29.05	T9
	ATOM	9453	CA	LYS	A	40	-61.374	116.910	-36.915	1.00	31.76	T9
	ATOM	9454	CB	LYS	A	40	-60.889	116.957	-38.364	1.00	28.39	T9
35	ATOM	9455	CG	LYS	A	40	-61.833	117.732	-39.272	1.00	30.19	T9
	ATOM	9456	CD	LYS	A	40	-61.547	117.455	-40.735	1.00	32.50	T9
	ATOM	9457	CE	LYS	A	40	-62.621	118.062	-41.624	1.00	30.08	T9
	ATOM	9458	NZ	LYS	A	40	-62.425	117.654	-43.047	1.00	30.67	T9
	ATOM	9459	C	LYS	A	40	-62.722	116.236	-36.849	1.00	32.25	T9
40	ATOM	9460	O	LYS	A	40	-62.925	115.171	-37.429	1.00	29.75	T9
	ATOM	9461	N	GLU	A	41	-63.634	116.853	-36.112	1.00	33.86	T9
	ATOM	9462	CA	GLU	A	41	-64.978	116.323	-35.978	1.00	28.86	T9
	ATOM	9463	CB	GLU	A	41	-65.731	116.561	-37.286	1.00	23.17	T9
	ATOM	9464	CG	GLU	A	41	-65.610	118.007	-37.733	1.00	33.16	T9
45	ATOM	9465	CD	GLU	A	41	-66.014	118.226	-39.181	1.00	20.49	T9
	ATOM	9466	OE1	GLU	A	41	-65.479	117.509	-40.077	1.00	29.15	T9
	ATOM	9467	OE2	GLU	A	41	-66.858	119.131	-39.428	1.00	24.99	T9
	ATOM	9468	C	GLU	A	41	-64.977	114.844	-35.608	1.00	29.93	T9
	ATOM	9469	O	GLU	A	41	-65.503	114.001	-36.330	1.00	27.95	T9
50	ATOM	9470	N	ASN	A	42	-64.355	114.546	-34.478	1.00	32.14	T9
	ATOM	9471	CA	ASN	A	42	-64.294	113.191	-33.968	1.00	33.10	T9
	ATOM	9472	CB	ASN	A	42	-65.694	112.682	-33.687	1.00	34.79	T9
	ATOM	9473	CG	ASN	A	42	-65.772	111.944	-32.390	1.00	30.51	T9
	ATOM	9474	OD1	ASN	A	42	-66.233	110.808	-32.338	1.00	31.13	T9
55	ATOM	9475	ND2	ASN	A	42	-65.318	112.586	-31.318	1.00	28.52	T9
	ATOM	9476	C	ASN	A	42	-63.580	112.185	-34.845	1.00	23.72	T9
	ATOM	9477	O	ASN	A	42	-63.797	110.980	-34.711	1.00	33.62	T9
	ATOM	9478	N	LYS	A	43	-62.730	112.679	-35.736	1.00	30.37	T9
	ATOM	9479	CA	LYS	A	43	-61.969	111.815	-36.623	1.00	31.24	T9
60	ATOM	9480	CB	LYS	A	43	-62.556	111.850	-38.034	1.00	29.00	T9
	ATOM	9481	CG	LYS	A	43	-63.909	111.179	-38.162	1.00	32.72	T9
	ATOM	9482	CD	LYS	A	43	-64.458	111.319	-39.568	1.00	30.48	T9
	ATOM	9483	CE	LYS	A	43	-64.873	112.754	-39.851	1.00	25.18	T9
	ATOM	9484	NZ	LYS	A	43	-65.451	112.925	-41.219	1.00	26.06	T9
65	ATOM	9485	C	LYS	A	43	-60.533	112.290	-36.658	1.00	33.42	T9
	ATOM	9486	O	LYS	A	43	-60.247	113.420	-36.293	1.00	31.90	T9

	ATOM	9487	N	ILE	A	44	-59.626	111.424	-37.084	1.00	28.76	T9
	ATOM	9488	CA	ILE	A	44	-58.231	111.804	-37.171	1.00	32.58	T9
	ATOM	9489	CB	ILE	A	44	-57.310	110.653	-36.777	1.00	31.99	T9
	ATOM	9490	CG2	ILE	A	44	-55.868	111.087	-36.895	1.00	25.05	T9
5	ATOM	9491	CG1	ILE	A	44	-57.608	110.218	-35.348	1.00	39.53	T9
	ATOM	9492	CD1	ILE	A	44	-56.762	109.080	-34.873	1.00	27.22	T9
	ATOM	9493	C	ILE	A	44	-57.949	112.197	-38.610	1.00	25.02	T9
	ATOM	9494	O	ILE	A	44	-58.177	111.421	-39.531	1.00	27.61	T9
	ATOM	9495	N	LEU	A	45	-57.455	113.413	-38.801	1.00	28.35	T9
10	ATOM	9496	CA	LEU	A	45	-57.162	113.917	-40.133	1.00	30.92	T9
	ATOM	9497	CB	LEU	A	45	-57.588	115.376	-40.225	1.00	34.49	T9
	ATOM	9498	CG	LEU	A	45	-57.258	116.035	-41.563	1.00	38.17	T9
	ATOM	9499	CD1	LEU	A	45	-58.149	115.466	-42.659	1.00	27.76	T9
	ATOM	9500	CD2	LEU	A	45	-57.456	117.527	-41.441	1.00	22.83	T9
15	ATOM	9501	C	LEU	A	45	-55.689	113.795	-40.515	1.00	19.49	T9
	ATOM	9502	O	LEU	A	45	-54.813	114.232	-39.772	1.00	34.57	T9
	ATOM	9503	N	VAL	A	46	-55.420	113.214	-41.680	1.00	30.20	T9
	ATOM	9504	CA	VAL	A	46	-54.052	113.047	-42.146	1.00	37.28	T9
	ATOM	9505	CB	VAL	A	46	-53.952	111.867	-43.103	1.00	35.28	T9
20	ATOM	9506	CG1	VAL	A	46	-52.537	111.720	-43.593	1.00	29.81	T9
	ATOM	9507	CG2	VAL	A	46	-54.393	110.612	-42.408	1.00	33.77	T9
	ATOM	9508	C	VAL	A	46	-53.566	114.299	-42.861	1.00	32.82	T9
	ATOM	9509	O	VAL	A	46	-54.146	114.705	-43.859	1.00	28.49	T9
	ATOM	9510	N	LYS	A	47	-52.497	114.908	-42.363	1.00	30.58	T9
25	ATOM	9511	CA	LYS	A	47	-51.974	116.125	-42.973	1.00	25.58	T9
	ATOM	9512	CB	LYS	A	47	-51.669	117.165	-41.893	1.00	28.31	T9
	ATOM	9513	CG	LYS	A	47	-52.882	117.857	-41.325	1.00	34.08	T9
	ATOM	9514	CD	LYS	A	47	-53.685	118.465	-42.442	1.00	31.30	T9
	ATOM	9515	CE	LYS	A	47	-54.741	119.428	-41.933	1.00	29.19	T9
30	ATOM	9516	NZ	LYS	A	47	-54.156	120.725	-41.466	1.00	30.86	T9
	ATOM	9517	C	LYS	A	47	-50.739	115.923	-43.839	1.00	29.14	T9
	ATOM	9518	O	LYS	A	47	-50.300	116.839	-44.515	1.00	29.50	T9
	ATOM	9519	N	GLU	A	48	-50.171	114.729	-43.805	1.00	26.57	T9
	ATOM	9520	CA	GLU	A	48	-48.984	114.409	-44.599	1.00	28.46	T9
35	ATOM	9521	CB	GLU	A	48	-47.715	114.531	-43.773	1.00	28.62	T9
	ATOM	9522	CG	GLU	A	48	-47.336	115.930	-43.383	1.00	29.64	T9
	ATOM	9523	CD	GLU	A	48	-46.148	115.940	-42.438	1.00	29.79	T9
	ATOM	9524	OE1	GLU	A	48	-45.205	115.132	-42.666	1.00	27.46	T9
	ATOM	9525	OE2	GLU	A	48	-46.157	116.753	-41.479	1.00	33.02	T9
40	ATOM	9526	C	GLU	A	48	-49.102	112.971	-45.027	1.00	29.14	T9
	ATOM	9527	O	GLU	A	48	-49.264	112.094	-44.184	1.00	34.93	T9
	ATOM	9528	N	THR	A	49	-49.013	112.708	-46.322	1.00	27.85	T9
	ATOM	9529	CA	THR	A	49	-49.134	111.332	-46.760	1.00	29.17	T9
	ATOM	9530	CB	THR	A	49	-49.284	111.239	-48.279	1.00	34.12	T9
45	ATOM	9531	OG1	THR	A	49	-47.997	111.087	-48.872	1.00	32.54	T9
	ATOM	9532	CG2	THR	A	49	-49.945	112.491	-48.818	1.00	32.51	T9
	ATOM	9533	C	THR	A	49	-47.904	110.547	-46.305	1.00	32.19	T9
	ATOM	9534	O	THR	A	49	-46.817	111.104	-46.160	1.00	26.66	T9
	ATOM	9535	N	GLY	A	50	-48.097	109.256	-46.055	1.00	26.03	T9
50	ATOM	9536	CA	GLY	A	50	-47.010	108.407	-45.609	1.00	34.58	T9
	ATOM	9537	C	GLY	A	50	-47.546	107.114	-45.027	1.00	28.29	T9
	ATOM	9538	O	GLY	A	50	-48.707	106.775	-45.239	1.00	37.23	T9
	ATOM	9539	N	TYR	A	51	-46.699	106.391	-44.301	1.00	31.20	T9
	ATOM	9540	CA	TYR	A	51	-47.093	105.137	-43.670	1.00	33.05	T9
55	ATOM	9541	CB	TYR	A	51	-46.000	104.078	-43.861	1.00	30.51	T9
	ATOM	9542	CG	TYR	A	51	-45.957	103.538	-45.263	1.00	26.33	T9
	ATOM	9543	CD1	TYR	A	51	-45.350	104.256	-46.291	1.00	30.89	T9
	ATOM	9544	CE1	TYR	A	51	-45.431	103.823	-47.618	1.00	41.01	T9
	ATOM	9545	CD2	TYR	A	51	-46.635	102.363	-45.589	1.00	26.82	T9
60	ATOM	9546	CE2	TYR	A	51	-46.727	101.918	-46.909	1.00	31.58	T9
	ATOM	9547	CZ	TYR	A	51	-46.130	102.656	-47.922	1.00	35.43	T9
	ATOM	9548	OH	TYR	A	51	-46.287	102.252	-49.234	1.00	23.09	T9
	ATOM	9549	C	TYR	A	51	-47.373	105.345	-42.182	1.00	24.75	T9
	ATOM	9550	O	TYR	A	51	-46.548	105.898	-41.451	1.00	30.02	T9
65	ATOM	9551	N	PHE	A	52	-48.543	104.899	-41.736	1.00	30.34	T9
	ATOM	9552	CA	PHE	A	52	-48.915	105.063	-40.345	1.00	34.17	T9

	ATOM	9553	CB	PHE	A	52	-50.096	106.023	-40.217	1.00	28.11	T9
	ATOM	9554	CG	PHE	A	52	-49.853	107.380	-40.796	1.00	28.64	T9
	ATOM	9555	CD1	PHE	A	52	-49.847	107.574	-42.167	1.00	32.14	T9
	ATOM	9556	CD2	PHE	A	52	-49.651	108.471	-39.966	1.00	32.39	T9
5	ATOM	9557	CE1	PHE	A	52	-49.644	108.839	-42.705	1.00	30.45	T9
	ATOM	9558	CE2	PHE	A	52	-49.448	109.734	-40.493	1.00	22.56	T9
	ATOM	9559	CZ	PHE	A	52	-49.444	109.919	-41.866	1.00	24.35	T9
	ATOM	9560	C	PHE	A	52	-49.290	103.765	-39.647	1.00	36.58	T9
	ATOM	9561	O	PHE	A	52	-49.811	102.838	-40.263	1.00	27.06	T9
10	ATOM	9562	N	PHE	A	53	-49.007	103.720	-38.350	1.00	28.98	T9
	ATOM	9563	CA	PHE	A	53	-49.344	102.590	-37.504	1.00	34.48	T9
	ATOM	9564	CB	PHE	A	53	-48.277	102.372	-36.442	1.00	31.55	T9
	ATOM	9565	CG	PHE	A	53	-48.662	101.370	-35.400	1.00	36.47	T9
	ATOM	9566	CD1	PHE	A	53	-48.802	100.031	-35.725	1.00	35.06	T9
15	ATOM	9567	CD2	PHE	A	53	-48.899	101.766	-34.088	1.00	33.34	T9
	ATOM	9568	CE1	PHE	A	53	-49.171	99.102	-34.759	1.00	27.59	T9
	ATOM	9569	CE2	PHE	A	53	-49.269	100.843	-33.120	1.00	38.75	T9
	ATOM	9570	CZ	PHE	A	53	-49.403	99.510	-33.459	1.00	33.60	T9
	ATOM	9571	C	PHE	A	53	-50.614	103.081	-36.844	1.00	30.83	T9
20	ATOM	9572	O	PHE	A	53	-50.619	104.136	-36.229	1.00	28.95	T9
	ATOM	9573	N	ILE	A	54	-51.695	102.328	-36.977	1.00	29.26	T9
	ATOM	9574	CA	ILE	A	54	-52.968	102.743	-36.400	1.00	26.88	T9
	ATOM	9575	CB	ILE	A	54	-53.992	102.947	-37.507	1.00	37.82	T9
	ATOM	9576	CG2	ILE	A	54	-55.243	103.581	-36.948	1.00	25.84	T9
25	ATOM	9577	CG1	ILE	A	54	-53.380	103.841	-38.583	1.00	34.36	T9
	ATOM	9578	CD1	ILE	A	54	-54.063	103.764	-39.898	1.00	24.51	T9
	ATOM	9579	C	ILE	A	54	-53.486	101.718	-35.409	1.00	30.44	T9
	ATOM	9580	O	ILE	A	54	-53.533	100.533	-35.717	1.00	33.96	T9
	ATOM	9581	N	TYR	A	55	-53.869	102.177	-34.220	1.00	27.96	T9
30	ATOM	9582	CA	TYR	A	55	-54.365	101.278	-33.188	1.00	30.65	T9
	ATOM	9583	CB	TYR	A	55	-53.305	101.098	-32.103	1.00	27.44	T9
	ATOM	9584	CG	TYR	A	55	-52.857	102.389	-31.470	1.00	26.56	T9
	ATOM	9585	CD1	TYR	A	55	-53.418	102.836	-30.285	1.00	33.89	T9
	ATOM	9586	CE1	TYR	A	55	-53.035	104.038	-29.717	1.00	33.41	T9
35	ATOM	9587	CD2	TYR	A	55	-51.894	103.179	-32.076	1.00	24.43	T9
	ATOM	9588	CE2	TYR	A	55	-51.503	104.390	-31.517	1.00	28.26	T9
	ATOM	9589	CZ	TYR	A	55	-52.079	104.812	-30.339	1.00	20.24	T9
	ATOM	9590	OH	TYR	A	55	-51.697	106.016	-29.796	1.00	33.02	T9
	ATOM	9591	C	TYR	A	55	-55.665	101.767	-32.570	1.00	26.13	T9
40	ATOM	9592	O	TYR	A	55	-56.112	102.871	-32.838	1.00	31.48	T9
	ATOM	9593	N	GLY	A	56	-56.276	100.927	-31.748	1.00	29.69	T9
	ATOM	9594	CA	GLY	A	56	-57.521	101.305	-31.116	1.00	30.12	T9
	ATOM	9595	C	GLY	A	56	-58.080	100.184	-30.271	1.00	28.15	T9
	ATOM	9596	O	GLY	A	56	-58.018	99.026	-30.664	1.00	33.55	T9
45	ATOM	9597	N	GLN	A	57	-58.612	100.537	-29.103	1.00	30.60	T9
	ATOM	9598	CA	GLN	A	57	-59.203	99.580	-28.176	1.00	28.94	T9
	ATOM	9599	CB	GLN	A	57	-58.270	99.318	-26.990	1.00	31.74	T9
	ATOM	9600	CG	GLN	A	57	-58.899	98.455	-25.902	1.00	33.28	T9
	ATOM	9601	CD	GLN	A	57	-57.945	98.082	-24.779	1.00	29.13	T9
50	ATOM	9602	OE1	GLN	A	57	-57.026	97.303	-24.969	1.00	27.71	T9
	ATOM	9603	NE2	GLN	A	57	-58.173	98.638	-23.598	1.00	34.56	T9
	ATOM	9604	C	GLN	A	57	-60.533	100.105	-27.661	1.00	20.21	T9
	ATOM	9605	O	GLN	A	57	-60.719	101.307	-27.514	1.00	26.87	T9
	ATOM	9606	N	VAL	A	58	-61.457	99.188	-27.396	1.00	30.52	T9
55	ATOM	9607	CA	VAL	A	58	-62.784	99.522	-26.887	1.00	27.36	T9
	ATOM	9608	CB	VAL	A	58	-63.849	99.500	-28.013	1.00	33.74	T9
	ATOM	9609	CG1	VAL	A	58	-65.238	99.576	-27.421	1.00	34.15	T9
	ATOM	9610	CG2	VAL	A	58	-63.630	100.660	-28.961	1.00	34.63	T9
	ATOM	9611	C	VAL	A	58	-63.167	98.485	-25.847	1.00	20.55	T9
60	ATOM	9612	O	VAL	A	58	-62.872	97.319	-26.023	1.00	32.79	T9
	ATOM	9613	N	LEU	A	59	-63.805	98.916	-24.761	1.00	34.45	T9
	ATOM	9614	CA	LEU	A	59	-64.244	97.997	-23.715	1.00	27.93	T9
	ATOM	9615	CB	LEU	A	59	-64.008	98.594	-22.329	1.00	28.66	T9
	ATOM	9616	CG	LEU	A	59	-63.872	97.601	-21.168	1.00	34.98	T9
65	ATOM	9617	CD1	LEU	A	59	-64.083	98.320	-19.866	1.00	29.60	T9
	ATOM	9618	CD2	LEU	A	59	-64.879	96.491	-21.285	1.00	36.21	T9

	ATOM	9619	C	LEU	A	59	-65.743	97.751	-23.905	1.00	28.68	T9
	ATOM	9620	O	LEU	A	59	-66.561	98.669	-23.736	1.00	27.32	T9
	ATOM	9621	N	TYR	A	60	-66.110	96.520	-24.253	1.00	27.19	T9
	ATOM	9622	CA	TYR	A	60	-67.518	96.202	-24.457	1.00	31.22	T9
5	ATOM	9623	CB	TYR	A	60	-67.664	95.180	-25.574	1.00	33.66	T9
	ATOM	9624	CG	TYR	A	60	-67.110	95.683	-26.857	1.00	30.43	T9
	ATOM	9625	CD1	TYR	A	60	-65.894	95.217	-27.344	1.00	36.03	T9
	ATOM	9626	CE1	TYR	A	60	-65.340	95.741	-28.511	1.00	26.98	T9
	ATOM	9627	CD2	TYR	A	60	-67.768	96.679	-27.562	1.00	33.29	T9
10	ATOM	9628	CE2	TYR	A	60	-67.234	97.214	-28.717	1.00	27.69	T9
	ATOM	9629	CZ	TYR	A	60	-66.020	96.744	-29.184	1.00	33.14	T9
	ATOM	9630	OH	TYR	A	60	-65.471	97.300	-30.308	1.00	33.36	T9
	ATOM	9631	C	TYR	A	60	-68.222	95.688	-23.208	1.00	32.60	T9
	ATOM	9632	O	TYR	A	60	-67.775	94.737	-22.567	1.00	28.53	T9
15	ATOM	9633	N	THR	A	61	-69.333	96.324	-22.870	1.00	32.18	T9
	ATOM	9634	CA	THR	A	61	-70.113	95.916	-21.716	1.00	27.23	T9
	ATOM	9635	CB	THR	A	61	-70.196	97.043	-20.699	1.00	31.36	T9
	ATOM	9636	OG1	THR	A	61	-70.685	98.229	-21.336	1.00	29.43	T9
	ATOM	9637	CG2	THR	A	61	-68.823	97.313	-20.115	1.00	30.29	T9
20	ATOM	9638	C	THR	A	61	-71.508	95.538	-22.183	1.00	28.04	T9
	ATOM	9639	O	THR	A	61	-72.438	95.429	-21.397	1.00	35.51	T9
	ATOM	9640	N	ASP	A	62	-71.629	95.338	-23.488	1.00	37.40	T9
	ATOM	9641	CA	ASP	A	62	-72.879	94.958	-24.127	1.00	27.54	T9
	ATOM	9642	CB	ASP	A	62	-72.853	95.442	-25.578	1.00	27.50	T9
25	ATOM	9643	CG	ASP	A	62	-74.208	95.378	-26.247	1.00	34.15	T9
	ATOM	9644	OD1	ASP	A	62	-74.609	96.409	-26.850	1.00	32.98	T9
	ATOM	9645	OD2	ASP	A	62	-74.855	94.300	-26.179	1.00	32.87	T9
	ATOM	9646	C	ASP	A	62	-72.966	93.435	-24.064	1.00	26.79	T9
	ATOM	9647	O	ASP	A	62	-71.943	92.760	-24.093	1.00	31.15	T9
30	ATOM	9648	N	LYS	A	63	-74.169	92.879	-23.976	1.00	25.78	T9
	ATOM	9649	CA	LYS	A	63	-74.272	91.425	-23.896	1.00	26.65	T9
	ATOM	9650	CB	LYS	A	63	-75.146	91.019	-22.707	1.00	25.55	T9
	ATOM	9651	CG	LYS	A	63	-76.576	91.548	-22.773	1.00	34.46	T9
	ATOM	9652	CD	LYS	A	63	-77.397	91.076	-21.568	1.00	30.36	T9
35	ATOM	9653	CE	LYS	A	63	-76.788	91.549	-20.222	1.00	25.26	T9
	ATOM	9654	NZ	LYS	A	63	-77.564	91.093	-19.013	1.00	33.00	T9
	ATOM	9655	C	LYS	A	63	-74.806	90.765	-25.153	1.00	33.25	T9
	ATOM	9656	O	LYS	A	63	-75.293	89.636	-25.097	1.00	25.54	T9
	ATOM	9657	N	THR	A	64	-74.697	91.436	-26.292	1.00	28.12	T9
40	ATOM	9658	CA	THR	A	64	-75.230	90.838	-27.500	1.00	27.17	T9
	ATOM	9659	CB	THR	A	64	-75.729	91.921	-28.495	1.00	34.69	T9
	ATOM	9660	OG1	THR	A	64	-74.710	92.897	-28.700	1.00	26.95	T9
	ATOM	9661	CG2	THR	A	64	-76.977	92.605	-27.955	1.00	29.76	T9
	ATOM	9662	C	THR	A	64	-74.333	89.839	-28.231	1.00	31.93	T9
45	ATOM	9663	O	THR	A	64	-73.978	90.037	-29.382	1.00	28.59	T9
	ATOM	9664	N	TYR	A	65	-73.986	88.758	-27.547	1.00	30.36	T9
	ATOM	9665	CA	TYR	A	65	-73.179	87.669	-28.104	1.00	29.71	T9
	ATOM	9666	CB	TYR	A	65	-74.080	86.750	-28.935	1.00	26.81	T9
	ATOM	9667	CG	TYR	A	65	-73.956	86.913	-30.428	1.00	25.97	T9
50	ATOM	9668	CD1	TYR	A	65	-73.056	86.139	-31.167	1.00	31.15	T9
	ATOM	9669	CE1	TYR	A	65	-72.934	86.295	-32.564	1.00	28.27	T9
	ATOM	9670	CD2	TYR	A	65	-74.730	87.843	-31.107	1.00	28.78	T9
	ATOM	9671	CE2	TYR	A	65	-74.621	88.012	-32.494	1.00	31.98	T9
	ATOM	9672	CZ	TYR	A	65	-73.726	87.239	-33.217	1.00	31.56	T9
55	ATOM	9673	OH	TYR	A	65	-73.629	87.424	-34.581	1.00	30.97	T9
	ATOM	9674	C	TYR	A	65	-71.911	87.981	-28.908	1.00	22.32	T9
	ATOM	9675	O	TYR	A	65	-71.009	87.140	-28.994	1.00	33.49	T9
	ATOM	9676	N	ALA	A	66	-71.825	89.163	-29.503	1.00	29.49	T9
	ATOM	9677	CA	ALA	A	66	-70.640	89.504	-30.277	1.00	31.65	T9
60	ATOM	9678	CB	ALA	A	66	-70.669	88.772	-31.600	1.00	29.05	T9
	ATOM	9679	C	ALA	A	66	-70.528	91.002	-30.508	1.00	27.98	T9
	ATOM	9680	O	ALA	A	66	-71.399	91.606	-31.123	1.00	35.78	T9
	ATOM	9681	N	MET	A	67	-69.454	91.597	-30.004	1.00	31.08	T9
	ATOM	9682	CA	MET	A	67	-69.227	93.020	-30.180	1.00	37.09	T9
65	ATOM	9683	CB	MET	A	67	-69.198	93.722	-28.832	1.00	24.01	T9
	ATOM	9684	CG	MET	A	67	-70.553	93.775	-28.145	1.00	26.41	T9

	ATOM	9685	SD	MET	A	67	-71.779	94.666	-29.147	1.00	31.12	T9
	ATOM	9686	CE	MET	A	67	-71.341	96.373	-28.791	1.00	30.56	T9
	ATOM	9687	C	MET	A	67	-67.909	93.220	-30.887	1.00	28.70	T9
	ATOM	9688	O	MET	A	67	-67.129	92.283	-31.018	1.00	30.90	T9
5	ATOM	9689	N	GLY	A	68	-67.663	94.440	-31.348	1.00	18.29	T9
	ATOM	9690	CA	GLY	A	68	-66.418	94.733	-32.041	1.00	29.77	T9
	ATOM	9691	C	GLY	A	68	-66.425	96.097	-32.705	1.00	33.15	T9
	ATOM	9692	O	GLY	A	68	-67.464	96.748	-32.773	1.00	23.19	T9
	ATOM	9693	N	HIS	A	69	-65.269	96.547	-33.180	1.00	32.58	T9
10	ATOM	9694	CA	HIS	A	69	-65.187	97.840	-33.847	1.00	36.28	T9
	ATOM	9695	CB	HIS	A	69	-64.626	98.919	-32.908	1.00	34.02	T9
	ATOM	9696	CG	HIS	A	69	-63.312	98.572	-32.275	1.00	35.94	T9
	ATOM	9697	CD2	HIS	A	69	-62.058	99.009	-32.535	1.00	20.04	T9
	ATOM	9698	ND1	HIS	A	69	-63.197	97.699	-31.216	1.00	37.18	T9
15	ATOM	9699	CE1	HIS	A	69	-61.933	97.614	-30.847	1.00	37.30	T9
	ATOM	9700	NE2	HIS	A	69	-61.220	98.399	-31.632	1.00	26.48	T9
	ATOM	9701	C	HIS	A	69	-64.366	97.790	-35.131	1.00	30.18	T9
	ATOM	9702	O	HIS	A	69	-63.676	96.815	-35.415	1.00	25.63	T9
	ATOM	9703	N	LEU	A	70	-64.457	98.858	-35.909	1.00	28.50	T9
20	ATOM	9704	CA	LEU	A	70	-63.746	98.957	-37.168	1.00	34.51	T9
	ATOM	9705	CB	LEU	A	70	-64.736	99.058	-38.318	1.00	28.57	T9
	ATOM	9706	CG	LEU	A	70	-65.993	98.198	-38.280	1.00	31.67	T9
	ATOM	9707	CD1	LEU	A	70	-66.910	98.637	-39.405	1.00	33.02	T9
	ATOM	9708	CD2	LEU	A	70	-65.630	96.735	-38.411	1.00	26.52	T9
25	ATOM	9709	C	LEU	A	70	-62.903	100.211	-37.198	1.00	28.87	T9
	ATOM	9710	O	LEU	A	70	-63.293	101.236	-36.656	1.00	37.95	T9
	ATOM	9711	N	ILE	A	71	-61.744	100.127	-37.832	1.00	36.93	T9
	ATOM	9712	CA	ILE	A	71	-60.892	101.296	-37.995	1.00	31.47	T9
	ATOM	9713	CB	ILE	A	71	-59.443	101.008	-37.598	1.00	34.39	T9
30	ATOM	9714	CG2	ILE	A	71	-58.548	102.132	-38.050	1.00	36.54	T9
	ATOM	9715	CG1	ILE	A	71	-59.351	100.865	-36.083	1.00	25.37	T9
	ATOM	9716	CD1	ILE	A	71	-57.958	100.553	-35.588	1.00	23.46	T9
	ATOM	9717	C	ILE	A	71	-60.992	101.583	-39.491	1.00	32.28	T9
	ATOM	9718	O	ILE	A	71	-60.477	100.833	-40.314	1.00	38.31	T9
35	ATOM	9719	N	GLN	A	72	-61.670	102.666	-39.845	1.00	24.46	T9
	ATOM	9720	CA	GLN	A	72	-61.871	102.977	-41.245	1.00	27.24	T9
	ATOM	9721	CB	GLN	A	72	-63.360	103.151	-41.497	1.00	30.39	T9
	ATOM	9722	CG	GLN	A	72	-64.175	102.066	-40.851	1.00	31.91	T9
	ATOM	9723	CD	GLN	A	72	-65.614	102.108	-41.277	1.00	28.73	T9
40	ATOM	9724	OE1	GLN	A	72	-66.275	103.139	-41.157	1.00	31.79	T9
	ATOM	9725	NE2	GLN	A	72	-66.117	100.984	-41.784	1.00	26.89	T9
	ATOM	9726	C	GLN	A	72	-61.117	104.173	-41.797	1.00	27.62	T9
	ATOM	9727	O	GLN	A	72	-60.727	105.082	-41.059	1.00	31.18	T9
	ATOM	9728	N	ARG	A	73	-60.942	104.155	-43.118	1.00	34.61	T9
45	ATOM	9729	CA	ARG	A	73	-60.240	105.200	-43.840	1.00	28.41	T9
	ATOM	9730	CB	ARG	A	73	-58.998	104.605	-44.484	1.00	27.89	T9
	ATOM	9731	CG	ARG	A	73	-58.197	105.593	-45.267	1.00	29.57	T9
	ATOM	9732	CD	ARG	A	73	-57.384	104.901	-46.320	1.00	27.78	T9
	ATOM	9733	NE	ARG	A	73	-56.620	105.859	-47.104	1.00	32.89	T9
50	ATOM	9734	CZ	ARG	A	73	-56.032	105.570	-48.256	1.00	31.52	T9
	ATOM	9735	NH1	ARG	A	73	-56.129	104.346	-48.760	1.00	24.09	T9
	ATOM	9736	NH2	ARG	A	73	-55.340	106.501	-48.893	1.00	31.03	T9
	ATOM	9737	C	ARG	A	73	-61.127	105.825	-44.921	1.00	29.98	T9
	ATOM	9738	O	ARG	A	73	-61.770	105.111	-45.680	1.00	34.42	T9
55	ATOM	9739	N	LYS	A	74	-61.170	107.155	-44.976	1.00	34.27	T9
	ATOM	9740	CA	LYS	A	74	-61.953	107.861	-45.988	1.00	36.28	T9
	ATOM	9741	CB	LYS	A	74	-62.755	108.999	-45.369	1.00	31.24	T9
	ATOM	9742	CG	LYS	A	74	-63.875	108.543	-44.454	1.00	33.41	T9
	ATOM	9743	CD	LYS	A	74	-64.603	109.727	-43.778	1.00	29.26	T9
60	ATOM	9744	CE	LYS	A	74	-65.715	109.237	-42.833	1.00	27.78	T9
	ATOM	9745	NZ	LYS	A	74	-66.482	110.352	-42.187	1.00	27.39	T9
	ATOM	9746	C	LYS	A	74	-60.992	108.451	-47.004	1.00	30.62	T9
	ATOM	9747	O	LYS	A	74	-60.391	109.498	-46.756	1.00	29.95	T9
	ATOM	9748	N	LYS	A	75	-60.844	107.782	-48.142	1.00	33.49	T9
65	ATOM	9749	CA	LYS	A	75	-59.944	108.238	-49.192	1.00	28.47	T9
	ATOM	9750	CB	LYS	A	75	-59.987	107.271	-50.375	1.00	34.22	T9

	ATOM	9751	CG	LYS	A	75	-59.582	105.845	-50.078	1.00	30.07	T9
	ATOM	9752	CD	LYS	A	75	-59.662	105.020	-51.354	1.00	32.53	T9
	ATOM	9753	CE	LYS	A	75	-59.257	103.570	-51.105	1.00	34.90	T9
	ATOM	9754	NZ	LYS	A	75	-59.419	102.651	-52.290	1.00	32.22	T9
5	ATOM	9755	C	LYS	A	75	-60.317	109.631	-49.692	1.00	32.06	T9
	ATOM	9756	O	LYS	A	75	-61.491	109.920	-49.897	1.00	33.78	T9
	ATOM	9757	N	VAL	A	76	-59.321	110.488	-49.895	1.00	30.41	T9
	ATOM	9758	CA	VAL	A	76	-59.571	111.836	-50.405	1.00	30.26	T9
	ATOM	9759	CB	VAL	A	76	-58.373	112.768	-50.233	1.00	25.56	T9
10	ATOM	9760	CG1	VAL	A	76	-58.812	114.188	-50.387	1.00	29.08	T9
	ATOM	9761	CG2	VAL	A	76	-57.726	112.546	-48.918	1.00	27.93	T9
	ATOM	9762	C	VAL	A	76	-59.759	111.713	-51.902	1.00	28.08	T9
	ATOM	9763	O	VAL	A	76	-60.612	112.373	-52.488	1.00	29.45	T9
	ATOM	9764	N	HIS	A	77	-58.926	110.873	-52.508	1.00	27.15	T9
15	ATOM	9765	CA	HIS	A	77	-58.962	110.636	-53.934	1.00	31.62	T9
	ATOM	9766	CB	HIS	A	77	-57.548	110.676	-54.490	1.00	38.98	T9
	ATOM	9767	CG	HIS	A	77	-56.841	111.972	-54.233	1.00	29.03	T9
	ATOM	9768	CD2	HIS	A	77	-55.523	112.286	-54.230	1.00	26.76	T9
	ATOM	9769	ND1	HIS	A	77	-57.516	113.155	-54.008	1.00	27.30	T9
20	ATOM	9770	CE1	HIS	A	77	-56.647	114.141	-53.883	1.00	34.17	T9
	ATOM	9771	NE2	HIS	A	77	-55.432	113.641	-54.016	1.00	30.16	T9
	ATOM	9772	C	HIS	A	77	-59.606	109.287	-54.195	1.00	29.51	T9
	ATOM	9773	O	HIS	A	77	-59.448	108.358	-53.406	1.00	21.37	T9
	ATOM	9774	N	VAL	A	78	-60.320	109.164	-55.310	1.00	31.08	T9
25	ATOM	9775	CA	VAL	A	78	-61.002	107.916	-55.576	1.00	28.40	T9
	ATOM	9776	CB	VAL	A	78	-62.521	108.119	-55.393	1.00	28.39	T9
	ATOM	9777	CG1	VAL	A	78	-63.254	106.799	-55.520	1.00	34.61	T9
	ATOM	9778	CG2	VAL	A	78	-62.784	108.697	-54.017	1.00	22.59	T9
	ATOM	9779	C	VAL	A	78	-60.722	107.137	-56.874	1.00	32.14	T9
30	ATOM	9780	O	VAL	A	78	-60.173	106.034	-56.810	1.00	22.98	T9
	ATOM	9781	N	PHE	A	79	-61.084	107.653	-58.041	1.00	28.08	T9
	ATOM	9782	CA	PHE	A	79	-60.836	106.895	-59.291	1.00	34.41	T9
	ATOM	9783	CB	PHE	A	79	-59.437	106.256	-59.326	1.00	22.40	T9
	ATOM	9784	CG	PHE	A	79	-58.319	107.209	-59.051	1.00	30.78	T9
35	ATOM	9785	CD1	PHE	A	79	-57.721	107.248	-57.796	1.00	29.95	T9
	ATOM	9786	CD2	PHE	A	79	-57.863	108.070	-60.039	1.00	35.19	T9
	ATOM	9787	CE1	PHE	A	79	-56.689	108.127	-57.528	1.00	33.50	T9
	ATOM	9788	CE2	PHE	A	79	-56.829	108.955	-59.784	1.00	28.70	T9
	ATOM	9789	CZ	PHE	A	79	-56.239	108.984	-58.525	1.00	32.03	T9
40	ATOM	9790	C	PHE	A	79	-61.824	105.753	-59.571	1.00	32.06	T9
	ATOM	9791	O	PHE	A	79	-61.963	104.818	-58.785	1.00	32.03	T9
	ATOM	9792	N	GLY	A	80	-62.479	105.825	-60.724	1.00	26.98	T9
	ATOM	9793	CA	GLY	A	80	-63.417	104.794	-61.128	1.00	30.45	T9
	ATOM	9794	C	GLY	A	80	-64.433	104.348	-60.107	1.00	28.37	T9
45	ATOM	9795	O	GLY	A	80	-65.107	105.161	-59.483	1.00	30.11	T9
	ATOM	9796	N	ASP	A	81	-64.537	103.036	-59.940	1.00	31.26	T9
	ATOM	9797	CA	ASP	A	81	-65.492	102.439	-59.015	1.00	28.78	T9
	ATOM	9798	CB	ASP	A	81	-66.148	101.210	-59.663	1.00	42.46	T9
	ATOM	9799	CG	ASP	A	81	-65.166	100.070	-59.907	1.00	30.79	T9
50	ATOM	9800	OD1	ASP	A	81	-63.949	100.306	-59.889	1.00	28.56	T9
	ATOM	9801	OD2	ASP	A	81	-65.607	98.929	-60.130	1.00	35.23	T9
	ATOM	9802	C	ASP	A	81	-64.906	102.060	-57.656	1.00	31.04	T9
	ATOM	9803	O	ASP	A	81	-65.366	101.119	-57.011	1.00	32.78	T9
	ATOM	9804	N	GLU	A	82	-63.888	102.789	-57.218	1.00	30.46	T9
55	ATOM	9805	CA	GLU	A	82	-63.295	102.515	-55.916	1.00	26.32	T9
	ATOM	9806	CB	GLU	A	82	-62.040	103.351	-55.686	1.00	27.82	T9
	ATOM	9807	CG	GLU	A	82	-60.781	102.870	-56.323	1.00	22.44	T9
	ATOM	9808	CD	GLU	A	82	-59.586	103.319	-55.523	1.00	24.50	T9
	ATOM	9809	OE1	GLU	A	82	-59.580	104.493	-55.097	1.00	31.18	T9
60	ATOM	9810	OE2	GLU	A	82	-58.654	102.510	-55.307	1.00	29.02	T9
	ATOM	9811	C	GLU	A	82	-64.297	102.961	-54.874	1.00	31.33	T9
	ATOM	9812	O	GLU	A	82	-65.061	103.884	-55.130	1.00	26.23	T9
	ATOM	9813	N	LEU	A	83	-64.307	102.321	-53.709	1.00	30.97	T9
	ATOM	9814	CA	LEU	A	83	-65.194	102.763	-52.641	1.00	30.33	T9
65	ATOM	9815	CB	LEU	A	83	-65.630	101.607	-51.750	1.00	30.12	T9
	ATOM	9816	CG	LEU	A	83	-66.573	100.574	-52.360	1.00	30.75	T9

	ATOM	9817	CD1	LEU	A	83	-65.869	99.851	-53.499	1.00	37.40	T9
	ATOM	9818	CD2	LEU	A	83	-66.993	99.584	-51.302	1.00	24.65	T9
	ATOM	9819	C	LEU	A	83	-64.284	103.681	-51.868	1.00	35.81	T9
	ATOM	9820	O	LEU	A	83	-63.154	103.312	-51.562	1.00	26.35	T9
5	ATOM	9821	N	SER	A	84	-64.756	104.883	-51.570	1.00	26.39	T9
	ATOM	9822	CA	SER	A	84	-63.931	105.846	-50.847	1.00	33.40	T9
	ATOM	9823	CB	SER	A	84	-64.452	107.267	-51.077	1.00	25.04	T9
	ATOM	9824	OG	SER	A	84	-65.851	107.329	-50.888	1.00	28.04	T9
	ATOM	9825	C	SER	A	84	-63.821	105.569	-49.356	1.00	33.40	T9
10	ATOM	9826	O	SER	A	84	-63.091	106.257	-48.642	1.00	32.49	T9
	ATOM	9827	N	LEU	A	85	-64.543	104.558	-48.886	1.00	30.71	T9
	ATOM	9828	CA	LEU	A	85	-64.503	104.185	-47.476	1.00	28.40	T9
	ATOM	9829	CB	LEU	A	85	-65.909	104.270	-46.875	1.00	23.54	T9
	ATOM	9830	CG	LEU	A	85	-66.089	104.316	-45.348	1.00	25.13	T9
15	ATOM	9831	CD1	LEU	A	85	-65.762	102.977	-44.717	1.00	33.91	T9
	ATOM	9832	CD2	LEU	A	85	-65.220	105.416	-44.767	1.00	24.26	T9
	ATOM	9833	C	LEU	A	85	-63.948	102.763	-47.352	1.00	31.31	T9
	ATOM	9834	O	LEU	A	85	-64.618	101.790	-47.678	1.00	30.35	T9
	ATOM	9835	N	VAL	A	86	-62.705	102.652	-46.904	1.00	30.23	T9
20	ATOM	9836	CA	VAL	A	86	-62.077	101.351	-46.737	1.00	29.49	T9
	ATOM	9837	CB	VAL	A	86	-60.658	101.316	-47.328	1.00	30.60	T9
	ATOM	9838	CG1	VAL	A	86	-60.011	99.967	-47.042	1.00	31.25	T9
	ATOM	9839	CG2	VAL	A	86	-60.705	101.574	-48.812	1.00	26.39	T9
	ATOM	9840	C	VAL	A	86	-61.949	101.059	-45.258	1.00	34.88	T9
25	ATOM	9841	O	VAL	A	86	-61.610	101.940	-44.476	1.00	30.20	T9
	ATOM	9842	N	THR	A	87	-62.219	99.824	-44.868	1.00	25.92	T9
	ATOM	9843	CA	THR	A	87	-62.099	99.485	-43.474	1.00	32.05	T9
	ATOM	9844	CB	THR	A	87	-63.377	98.784	-42.933	1.00	30.17	T9
	ATOM	9845	OG1	THR	A	87	-63.054	97.463	-42.506	1.00	29.15	T9
30	ATOM	9846	CG2	THR	A	87	-64.465	98.739	-43.999	1.00	26.12	T9
	ATOM	9847	C	THR	A	87	-60.876	98.606	-43.305	1.00	31.02	T9
	ATOM	9848	O	THR	A	87	-60.782	97.537	-43.876	1.00	28.87	T9
	ATOM	9849	N	LEU	A	88	-59.922	99.101	-42.532	1.00	25.92	T9
	ATOM	9850	CA	LEU	A	88	-58.674	98.408	-42.242	1.00	35.21	T9
35	ATOM	9851	CB	LEU	A	88	-57.539	99.425	-42.104	1.00	30.20	T9
	ATOM	9852	CG	LEU	A	88	-57.145	100.460	-43.168	1.00	31.82	T9
	ATOM	9853	CD1	LEU	A	88	-58.319	100.855	-44.009	1.00	33.27	T9
	ATOM	9854	CD2	LEU	A	88	-56.579	101.681	-42.479	1.00	35.37	T9
	ATOM	9855	C	LEU	A	88	-58.842	97.731	-40.890	1.00	35.16	T9
40	ATOM	9856	O	LEU	A	88	-59.526	98.248	-40.002	1.00	37.36	T9
	ATOM	9857	N	PHE	A	89	-58.253	96.565	-40.711	1.00	29.45	T9
	ATOM	9858	CA	PHE	A	89	-58.326	95.939	-39.386	1.00	30.03	T9
	ATOM	9859	CB	PHE	A	89	-57.705	96.902	-38.363	1.00	24.78	T9
	ATOM	9860	CG	PHE	A	89	-56.448	97.553	-38.865	1.00	31.39	T9
45	ATOM	9861	CD1	PHE	A	89	-56.137	98.860	-38.518	1.00	27.45	T9
	ATOM	9862	CD2	PHE	A	89	-55.623	96.880	-39.784	1.00	31.42	T9
	ATOM	9863	CE1	PHE	A	89	-55.028	99.494	-39.092	1.00	25.56	T9
	ATOM	9864	CE2	PHE	A	89	-54.516	97.501	-40.360	1.00	32.57	T9
	ATOM	9865	CZ	PHE	A	89	-54.217	98.807	-40.021	1.00	30.36	T9
50	ATOM	9866	C	PHE	A	89	-59.667	95.412	-38.862	1.00	34.47	T9
	ATOM	9867	O	PHE	A	89	-60.116	94.361	-39.306	1.00	32.30	T9
	ATOM	9868	N	ARG	A	90	-60.293	96.091	-37.904	1.00	28.56	T9
	ATOM	9869	CA	ARG	A	90	-61.554	95.566	-37.358	1.00	36.95	T9
	ATOM	9870	CB	ARG	A	90	-62.538	95.295	-38.494	1.00	28.05	T9
55	ATOM	9871	CG	ARG	A	90	-63.545	94.199	-38.198	1.00	33.70	T9
	ATOM	9872	CD	ARG	A	90	-64.206	93.702	-39.466	1.00	32.27	T9
	ATOM	9873	NE	ARG	A	90	-64.995	92.504	-39.198	1.00	31.94	T9
	ATOM	9874	CZ	ARG	A	90	-65.118	91.482	-40.039	1.00	24.59	T9
	ATOM	9875	NH1	ARG	A	90	-64.503	91.504	-41.219	1.00	37.78	T9
60	ATOM	9876	NH2	ARG	A	90	-65.842	90.425	-39.689	1.00	31.80	T9
	ATOM	9877	C	ARG	A	90	-61.411	94.269	-36.528	1.00	30.18	T9
	ATOM	9878	O	ARG	A	90	-60.841	93.288	-36.989	1.00	26.12	T9
	ATOM	9879	N	CYS	A	91	-61.966	94.264	-35.320	1.00	37.52	T9
	ATOM	9880	CA	CYS	A	91	-61.908	93.092	-34.440	1.00	39.42	T9
65	ATOM	9881	CB	CYS	A	91	-60.951	93.367	-33.284	1.00	31.58	T9
	ATOM	9882	SG	CYS	A	91	-61.358	94.866	-32.372	1.00	30.56	T9

	ATOM	9883	C	CYS	A	91	-63.287	92.691	-33.888	1.00	34.61	T9
	ATOM	9884	O	CYS	A	91	-64.244	93.457	-33.972	1.00	32.46	T9
	ATOM	9885	N	ILE	A	92	-63.377	91.493	-33.315	1.00	30.06	T9
	ATOM	9886	CA	ILE	A	92	-64.637	90.972	-32.775	1.00	21.85	T9
5	ATOM	9887	CB	ILE	A	92	-65.271	89.952	-33.739	1.00	27.47	T9
	ATOM	9888	CG2	ILE	A	92	-66.602	89.487	-33.209	1.00	29.46	T9
	ATOM	9889	CG1	ILE	A	92	-65.472	90.587	-35.111	1.00	32.28	T9
	ATOM	9890	CD1	ILE	A	92	-65.793	89.580	-36.203	1.00	33.11	T9
	ATOM	9891	C	ILE	A	92	-64.407	90.253	-31.458	1.00	33.96	T9
10	ATOM	9892	O	ILE	A	92	-63.326	89.760	-31.211	1.00	38.51	T9
	ATOM	9893	N	GLN	A	93	-65.428	90.191	-30.614	1.00	26.80	T9
	ATOM	9894	CA	GLN	A	93	-65.318	89.503	-29.330	1.00	34.57	T9
	ATOM	9895	CB	GLN	A	93	-64.760	90.437	-28.250	1.00	24.28	T9
	ATOM	9896	CG	GLN	A	93	-63.241	90.496	-28.185	1.00	33.62	T9
15	ATOM	9897	CD	GLN	A	93	-62.661	89.585	-27.118	1.00	31.97	T9
	ATOM	9898	OE1	GLN	A	93	-62.604	88.366	-27.273	1.00	28.43	T9
	ATOM	9899	NE2	GLN	A	93	-62.231	90.180	-26.013	1.00	28.56	T9
	ATOM	9900	C	GLN	A	93	-66.667	88.964	-28.878	1.00	35.27	T9
	ATOM	9901	O	GLN	A	93	-67.626	89.720	-28.719	1.00	30.63	T9
20	ATOM	9902	N	ASN	A	94	-66.742	87.652	-28.689	1.00	32.63	T9
	ATOM	9903	CA	ASN	A	94	-67.979	87.044	-28.238	1.00	33.39	T9
	ATOM	9904	CB	ASN	A	94	-67.841	85.515	-28.143	1.00	28.05	T9
	ATOM	9905	CG	ASN	A	94	-68.005	84.822	-29.493	1.00	27.18	T9
	ATOM	9906	OD1	ASN	A	94	-69.004	85.013	-30.178	1.00	32.28	T9
25	ATOM	9907	ND2	ASN	A	94	-67.032	84.009	-29.869	1.00	28.23	T9
	ATOM	9908	C	ASN	A	94	-68.246	87.636	-26.865	1.00	31.56	T9
	ATOM	9909	O	ASN	A	94	-67.308	87.927	-26.119	1.00	37.47	T9
	ATOM	9910	N	MET	A	95	-69.520	87.831	-26.542	1.00	30.97	T9
	ATOM	9911	CA	MET	A	95	-69.896	88.388	-25.252	1.00	30.63	T9
30	ATOM	9912	CB	MET	A	95	-70.773	89.630	-25.445	1.00	39.15	T9
	ATOM	9913	CG	MET	A	95	-70.103	90.760	-26.199	1.00	31.70	T9
	ATOM	9914	SD	MET	A	95	-68.485	91.230	-25.514	1.00	25.49	T9
	ATOM	9915	CE	MET	A	95	-68.957	92.085	-23.998	1.00	36.82	T9
	ATOM	9916	C	MET	A	95	-70.662	87.365	-24.417	1.00	31.20	T9
35	ATOM	9917	O	MET	A	95	-71.330	86.479	-24.963	1.00	31.51	T9
	ATOM	9918	N	PRO	A	96	-70.560	87.467	-23.077	1.00	27.70	T9
	ATOM	9919	CD	PRO	A	96	-69.684	88.377	-22.326	1.00	34.07	T9
	ATOM	9920	CA	PRO	A	96	-71.251	86.562	-22.158	1.00	28.61	T9
	ATOM	9921	CB	PRO	A	96	-70.556	86.800	-20.819	1.00	37.05	T9
40	ATOM	9922	CG	PRO	A	96	-69.288	87.502	-21.179	1.00	32.99	T9
	ATOM	9923	C	PRO	A	96	-72.683	87.058	-22.106	1.00	35.73	T9
	ATOM	9924	O	PRO	A	96	-73.080	87.933	-22.877	1.00	29.09	T9
	ATOM	9925	N	GLU	A	97	-73.455	86.535	-21.171	1.00	31.26	T9
	ATOM	9926	CA	GLU	A	97	-74.834	86.953	-21.059	1.00	29.24	T9
45	ATOM	9927	CB	GLU	A	97	-75.740	85.736	-21.150	1.00	30.21	T9
	ATOM	9928	CG	GLU	A	97	-77.177	86.092	-21.412	1.00	31.42	T9
	ATOM	9929	CD	GLU	A	97	-77.776	85.190	-22.460	1.00	23.42	T9
	ATOM	9930	OE1	GLU	A	97	-77.817	83.954	-22.214	1.00	35.56	T9
	ATOM	9931	OE2	GLU	A	97	-78.195	85.714	-23.529	1.00	28.50	T9
50	ATOM	9932	C	GLU	A	97	-75.061	87.665	-19.743	1.00	34.70	T9
	ATOM	9933	O	GLU	A	97	-75.992	88.465	-19.604	1.00	25.51	T9
	ATOM	9934	N	THR	A	98	-74.187	87.382	-18.785	1.00	27.93	T9
	ATOM	9935	CA	THR	A	98	-74.299	87.960	-17.462	1.00	29.68	T9
	ATOM	9936	CB	THR	A	98	-73.872	86.954	-16.420	1.00	32.37	T9
55	ATOM	9937	OG1	THR	A	98	-72.483	86.651	-16.622	1.00	30.60	T9
	ATOM	9938	CG2	THR	A	98	-74.702	85.676	-16.551	1.00	31.10	T9
	ATOM	9939	C	THR	A	98	-73.511	89.245	-17.225	1.00	35.25	T9
	ATOM	9940	O	THR	A	98	-74.098	90.320	-17.129	1.00	32.03	T9
	ATOM	9941	N	LEU	A	99	-72.193	89.160	-17.135	1.00	34.56	T9
60	ATOM	9942	CA	LEU	A	99	-71.441	90.367	-16.849	1.00	29.26	T9
	ATOM	9943	CB	LEU	A	99	-70.707	90.188	-15.522	1.00	38.26	T9
	ATOM	9944	CG	LEU	A	99	-71.673	89.988	-14.347	1.00	27.34	T9
	ATOM	9945	CD1	LEU	A	99	-70.908	89.657	-13.082	1.00	37.97	T9
	ATOM	9946	CD2	LEU	A	99	-72.493	91.250	-14.156	1.00	33.36	T9
65	ATOM	9947	C	LEU	A	99	-70.480	90.806	-17.939	1.00	29.90	T9
	ATOM	9948	O	LEU	A	99	-69.262	90.709	-17.788	1.00	34.71	T9

	ATOM	9949	N	PRO	A	100	-71.023	91.325	-19.050	1.00	29.49	T9
	ATOM	9950	CD	PRO	A	100	-72.456	91.575	-19.274	1.00	28.06	T9
	ATOM	9951	CA	PRO	A	100	-70.240	91.793	-20.192	1.00	27.14	T9
	ATOM	9952	CB	PRO	A	100	-71.261	92.569	-21.010	1.00	35.36	T9
5	ATOM	9953	CG	PRO	A	100	-72.511	91.791	-20.769	1.00	19.96	T9
	ATOM	9954	C	PRO	A	100	-69.047	92.650	-19.794	1.00	30.62	T9
	ATOM	9955	O	PRO	A	100	-69.191	93.671	-19.128	1.00	34.14	T9
	ATOM	9956	N	ASN	A	101	-67.866	92.221	-20.218	1.00	26.57	T9
	ATOM	9957	CA	ASN	A	101	-66.631	92.923	-19.933	1.00	31.77	T9
10	ATOM	9958	CB	ASN	A	101	-66.213	92.687	-18.501	1.00	35.14	T9
	ATOM	9959	CG	ASN	A	101	-66.826	93.662	-17.573	1.00	31.48	T9
	ATOM	9960	OD1	ASN	A	101	-66.497	94.844	-17.607	1.00	37.83	T9
	ATOM	9961	ND2	ASN	A	101	-67.740	93.190	-16.734	1.00	27.10	T9
	ATOM	9962	C	ASN	A	101	-65.543	92.396	-20.818	1.00	30.11	T9
15	ATOM	9963	O	ASN	A	101	-64.770	91.539	-20.377	1.00	29.41	T9
	ATOM	9964	N	ASN	A	102	-65.447	92.882	-22.055	1.00	32.35	T9
	ATOM	9965	CA	ASN	A	102	-64.394	92.340	-22.879	1.00	28.43	T9
	ATOM	9966	CB	ASN	A	102	-64.987	91.513	-24.018	1.00	32.44	T9
	ATOM	9967	CG	ASN	A	102	-65.293	90.088	-23.579	1.00	38.48	T9
20	ATOM	9968	OD1	ASN	A	102	-64.476	89.443	-22.914	1.00	28.15	T9
	ATOM	9969	ND2	ASN	A	102	-66.469	89.591	-23.941	1.00	30.24	T9
	ATOM	9970	C	ASN	A	102	-63.244	93.199	-23.366	1.00	28.58	T9
	ATOM	9971	O	ASN	A	102	-62.094	92.768	-23.235	1.00	32.14	T9
	ATOM	9972	N	SER	A	103	-63.472	94.388	-23.906	1.00	31.32	T9
25	ATOM	9973	CA	SER	A	103	-62.288	95.143	-24.349	1.00	31.96	T9
	ATOM	9974	CB	SER	A	103	-61.361	95.427	-23.149	1.00	26.91	T9
	ATOM	9975	OG	SER	A	103	-60.006	95.125	-23.438	1.00	32.56	T9
	ATOM	9976	C	SER	A	103	-61.503	94.382	-25.435	1.00	32.63	T9
	ATOM	9977	O	SER	A	103	-61.162	93.207	-25.294	1.00	30.84	T9
30	ATOM	9978	N	CYS	A	104	-61.202	95.064	-26.526	1.00	27.30	T9
	ATOM	9979	CA	CYS	A	104	-60.487	94.419	-27.601	1.00	32.19	T9
	ATOM	9980	CB	CYS	A	104	-61.496	93.838	-28.583	1.00	32.02	T9
	ATOM	9981	SG	CYS	A	104	-60.816	92.731	-29.803	1.00	38.77	T9
	ATOM	9982	C	CYS	A	104	-59.585	95.421	-28.291	1.00	26.21	T9
35	ATOM	9983	O	CYS	A	104	-60.031	96.487	-28.691	1.00	27.66	T9
	ATOM	9984	N	TYR	A	105	-58.311	95.072	-28.412	1.00	30.04	T9
	ATOM	9985	CA	TYR	A	105	-57.324	95.924	-29.054	1.00	30.51	T9
	ATOM	9986	CB	TYR	A	105	-56.070	95.998	-28.193	1.00	26.73	T9
	ATOM	9987	CG	TYR	A	105	-54.924	96.771	-28.798	1.00	28.79	T9
40	ATOM	9988	CD1	TYR	A	105	-54.742	98.114	-28.513	1.00	31.78	T9
	ATOM	9989	CE1	TYR	A	105	-53.675	98.822	-29.051	1.00	36.89	T9
	ATOM	9990	CD2	TYR	A	105	-54.010	96.153	-29.642	1.00	27.98	T9
	ATOM	9991	CE2	TYR	A	105	-52.945	96.855	-30.185	1.00	28.36	T9
	ATOM	9992	CZ	TYR	A	105	-52.785	98.185	-29.883	1.00	27.12	T9
45	ATOM	9993	OH	TYR	A	105	-51.725	98.875	-30.397	1.00	29.40	T9
	ATOM	9994	C	TYR	A	105	-56.960	95.345	-30.408	1.00	31.98	T9
	ATOM	9995	O	TYR	A	105	-56.955	94.137	-30.593	1.00	27.25	T9
	ATOM	9996	N	SER	A	106	-56.661	96.213	-31.358	1.00	34.88	T9
	ATOM	9997	CA	SER	A	106	-56.267	95.772	-32.680	1.00	24.08	T9
50	ATOM	9998	CB	SER	A	106	-57.491	95.378	-33.508	1.00	38.25	T9
	ATOM	9999	OG	SER	A	106	-57.120	94.714	-34.707	1.00	25.09	T9
	ATOM	10000	C	SER	A	106	-55.545	96.939	-33.321	1.00	32.60	T9
	ATOM	10001	O	SER	A	106	-55.914	98.089	-33.090	1.00	27.72	T9
	ATOM	10002	N	ALA	A	107	-54.509	96.637	-34.104	1.00	29.80	T9
55	ATOM	10003	CA	ALA	A	107	-53.714	97.658	-34.777	1.00	30.76	T9
	ATOM	10004	CB	ALA	A	107	-52.670	98.203	-33.823	1.00	35.17	T9
	ATOM	10005	C	ALA	A	107	-53.038	97.117	-36.029	1.00	34.22	T9
	ATOM	10006	O	ALA	A	107	-52.942	95.913	-36.227	1.00	32.23	T9
	ATOM	10007	N	GLY	A	108	-52.566	98.023	-36.873	1.00	24.00	T9
60	ATOM	10008	CA	GLY	A	108	-51.889	97.624	-38.091	1.00	31.83	T9
	ATOM	10009	C	GLY	A	108	-51.266	98.817	-38.790	1.00	33.97	T9
	ATOM	10010	O	GLY	A	108	-51.375	99.944	-38.316	1.00	30.97	T9
	ATOM	10011	N	ILE	A	109	-50.610	98.574	-39.917	1.00	31.44	T9
	ATOM	10012	CA	ILE	A	109	-49.970	99.640	-40.677	1.00	30.80	T9
65	ATOM	10013	CB	ILE	A	109	-48.510	99.297	-40.993	1.00	28.58	T9
	ATOM	10014	CG2	ILE	A	109	-47.883	100.404	-41.814	1.00	34.25	T9

	ATOM	10015	CG1	ILE	A	109	-47.732	99.086	-39.696	1.00	24.19	T9
	ATOM	10016	CD1	ILE	A	109	-46.349	98.540	-39.916	1.00	30.88	T9
	ATOM	10017	C	ILE	A	109	-50.704	99.834	-41.993	1.00	29.02	T9
	ATOM	10018	O	ILE	A	109	-51.137	98.869	-42.614	1.00	34.90	T9
5	ATOM	10019	N	ALA	A	110	-50.843	101.082	-42.417	1.00	27.82	T9
	ATOM	10020	CA	ALA	A	110	-51.530	101.385	-43.661	1.00	25.18	T9
	ATOM	10021	CB	ALA	A	110	-53.005	101.597	-43.398	1.00	35.07	T9
	ATOM	10022	C	ALA	A	110	-50.929	102.641	-44.268	1.00	30.49	T9
	ATOM	10023	O	ALA	A	110	-50.357	103.467	-43.555	1.00	29.63	T9
10	ATOM	10024	N	LYS	A	111	-51.040	102.783	-45.585	1.00	29.13	T9
	ATOM	10025	CA	LYS	A	111	-50.517	103.970	-46.231	1.00	30.00	T9
	ATOM	10026	CB	LYS	A	111	-49.944	103.640	-47.602	1.00	26.01	T9
	ATOM	10027	CG	LYS	A	111	-49.334	104.863	-48.271	1.00	33.91	T9
	ATOM	10028	CD	LYS	A	111	-48.712	104.533	-49.620	1.00	37.00	T9
15	ATOM	10029	CE	LYS	A	111	-48.085	105.777	-50.262	1.00	30.28	T9
	ATOM	10030	NZ	LYS	A	111	-47.432	105.471	-51.585	1.00	30.61	T9
	ATOM	10031	C	LYS	A	111	-51.673	104.941	-46.372	1.00	30.13	T9
	ATOM	10032	O	LYS	A	111	-52.748	104.556	-46.814	1.00	34.78	T9
	ATOM	10033	N	LEU	A	112	-51.447	106.194	-45.987	1.00	35.29	T9
20	ATOM	10034	CA	LEU	A	112	-52.476	107.229	-46.041	1.00	36.89	T9
	ATOM	10035	CB	LEU	A	112	-52.866	107.639	-44.625	1.00	34.93	T9
	ATOM	10036	CG	LEU	A	112	-53.192	106.517	-43.648	1.00	23.16	T9
	ATOM	10037	CD1	LEU	A	112	-53.262	107.062	-42.247	1.00	26.47	T9
	ATOM	10038	CD2	LEU	A	112	-54.496	105.867	-44.048	1.00	28.57	T9
25	ATOM	10039	C	LEU	A	112	-51.969	108.462	-46.775	1.00	32.21	T9
	ATOM	10040	O	LEU	A	112	-50.765	108.643	-46.914	1.00	33.94	T9
	ATOM	10041	N	GLU	A	113	-52.891	109.314	-47.223	1.00	26.60	T9
	ATOM	10042	CA	GLU	A	113	-52.535	110.537	-47.938	1.00	23.82	T9
	ATOM	10043	CB	GLU	A	113	-52.983	110.483	-49.389	1.00	26.47	T9
30	ATOM	10044	CG	GLU	A	113	-52.815	109.178	-50.091	1.00	26.73	T9
	ATOM	10045	CD	GLU	A	113	-52.831	109.362	-51.605	1.00	29.37	T9
	ATOM	10046	OE1	GLU	A	113	-53.703	110.119	-52.122	1.00	30.10	T9
	ATOM	10047	OE2	GLU	A	113	-51.961	108.747	-52.281	1.00	32.65	T9
	ATOM	10048	C	GLU	A	113	-53.182	111.772	-47.345	1.00	29.38	T9
35	ATOM	10049	O	GLU	A	113	-54.214	111.682	-46.685	1.00	29.94	T9
	ATOM	10050	N	GLU	A	114	-52.581	112.929	-47.616	1.00	28.39	T9
	ATOM	10051	CA	GLU	A	114	-53.114	114.208	-47.156	1.00	32.22	T9
	ATOM	10052	CB	GLU	A	114	-52.546	115.356	-47.961	1.00	32.08	T9
	ATOM	10053	CG	GLU	A	114	-51.278	115.945	-47.466	1.00	27.52	T9
40	ATOM	10054	CD	GLU	A	114	-51.079	117.329	-48.052	1.00	30.82	T9
	ATOM	10055	OE1	GLU	A	114	-51.909	118.229	-47.760	1.00	30.34	T9
	ATOM	10056	OE2	GLU	A	114	-50.101	117.507	-48.814	1.00	28.31	T9
	ATOM	10057	C	GLU	A	114	-54.604	114.246	-47.393	1.00	27.51	T9
	ATOM	10058	O	GLU	A	114	-55.046	114.053	-48.518	1.00	20.40	T9
45	ATOM	10059	N	GLY	A	115	-55.382	114.508	-46.357	1.00	31.15	T9
	ATOM	10060	CA	GLY	A	115	-56.812	114.590	-46.551	1.00	30.83	T9
	ATOM	10061	C	GLY	A	115	-57.570	113.356	-46.144	1.00	32.96	T9
	ATOM	10062	O	GLY	A	115	-58.787	113.417	-45.967	1.00	30.83	T9
	ATOM	10063	N	ASP	A	116	-56.885	112.225	-46.021	1.00	25.06	T9
50	ATOM	10064	CA	ASP	A	116	-57.583	111.017	-45.601	1.00	22.83	T9
	ATOM	10065	CB	ASP	A	116	-56.674	109.785	-45.671	1.00	27.23	T9
	ATOM	10066	CG	ASP	A	116	-56.359	109.353	-47.096	1.00	26.11	T9
	ATOM	10067	OD1	ASP	A	116	-57.173	109.628	-48.010	1.00	27.90	T9
	ATOM	10068	OD2	ASP	A	116	-55.302	108.709	-47.300	1.00	32.72	T9
55	ATOM	10069	C	ASP	A	116	-58.026	111.223	-44.153	1.00	26.70	T9
	ATOM	10070	O	ASP	A	116	-57.411	111.990	-43.401	1.00	34.85	T9
	ATOM	10071	N	GLU	A	117	-59.099	110.553	-43.758	1.00	26.01	T9
	ATOM	10072	CA	GLU	A	117	-59.566	110.667	-42.389	1.00	31.13	T9
	ATOM	10073	CB	GLU	A	117	-60.910	111.380	-42.335	1.00	26.88	T9
60	ATOM	10074	CG	GLU	A	117	-60.914	112.721	-43.009	1.00	32.45	T9
	ATOM	10075	CD	GLU	A	117	-62.246	113.425	-42.866	1.00	28.69	T9
	ATOM	10076	OE1	GLU	A	117	-63.292	112.729	-42.908	1.00	28.26	T9
	ATOM	10077	OE2	GLU	A	117	-62.247	114.671	-42.724	1.00	30.39	T9
	ATOM	10078	C	GLU	A	117	-59.710	109.267	-41.823	1.00	35.25	T9
65	ATOM	10079	O	GLU	A	117	-60.123	108.352	-42.529	1.00	30.93	T9
	ATOM	10080	N	LEU	A	118	-59.346	109.107	-40.555	1.00	36.08	T9

	ATOM	10081	CA	LEU A 118	-59.459	107.829	-39.877	1.00	30.33	T9
	ATOM	10082	CB	LEU A 118	-58.161	107.483	-39.162	1.00	27.53	T9
	ATOM	10083	CG	LEU A 118	-56.940	107.234	-40.031	1.00	35.85	T9
	ATOM	10084	CD1	LEU A 118	-55.775	106.839	-39.157	1.00	26.77	T9
5	ATOM	10085	CD2	LEU A 118	-57.246	106.138	-41.023	1.00	22.58	T9
	ATOM	10086	C	LEU A 118	-60.568	107.940	-38.852	1.00	27.04	T9
	ATOM	10087	O	LEU A 118	-60.765	108.992	-38.261	1.00	28.77	T9
	ATOM	10088	N	GLN A 119	-61.298	106.851	-38.648	1.00	26.36	T9
	ATOM	10089	CA	GLN A 119	-62.374	106.834	-37.666	1.00	26.27	T9
10	ATOM	10090	CB	GLN A 119	-63.659	107.331	-38.300	1.00	27.74	T9
	ATOM	10091	CG	GLN A 119	-64.155	106.465	-39.436	1.00	34.21	T9
	ATOM	10092	CD	GLN A 119	-65.395	107.035	-40.102	1.00	30.46	T9
	ATOM	10093	OE1	GLN A 119	-66.116	106.325	-40.798	1.00	28.88	T9
	ATOM	10094	NE2	GLN A 119	-65.642	108.324	-39.898	1.00	35.84	T9
15	ATOM	10095	C	GLN A 119	-62.580	105.437	-37.089	1.00	34.55	T9
	ATOM	10096	O	GLN A 119	-62.251	104.440	-37.725	1.00	26.84	T9
	ATOM	10097	N	LEU A 120	-63.119	105.385	-35.876	1.00	30.99	T9
	ATOM	10098	CA	LEU A 120	-63.377	104.139	-35.154	1.00	34.87	T9
	ATOM	10099	CB	LEU A 120	-62.817	104.264	-33.732	1.00	39.11	T9
20	ATOM	10100	CG	LEU A 120	-62.644	103.103	-32.749	1.00	28.00	T9
	ATOM	10101	CD1	LEU A 120	-63.882	102.237	-32.722	1.00	36.79	T9
	ATOM	10102	CD2	LEU A 120	-61.438	102.292	-33.139	1.00	25.01	T9
	ATOM	10103	C	LEU A 120	-64.888	103.960	-35.098	1.00	27.44	T9
	ATOM	10104	O	LEU A 120	-65.591	104.808	-34.554	1.00	37.91	T9
25	ATOM	10105	N	ALA A 121	-65.396	102.862	-35.643	1.00	29.10	T9
	ATOM	10106	CA	ALA A 121	-66.837	102.646	-35.650	1.00	31.17	T9
	ATOM	10107	CB	ALA A 121	-67.370	102.837	-37.057	1.00	29.70	T9
	ATOM	10108	C	ALA A 121	-67.320	101.302	-35.098	1.00	26.68	T9
	ATOM	10109	O	ALA A 121	-66.710	100.255	-35.317	1.00	26.76	T9
30	ATOM	10110	N	ILE A 122	-68.432	101.347	-34.373	1.00	32.47	T9
	ATOM	10111	CA	ILE A 122	-69.029	100.146	-33.813	1.00	31.82	T9
	ATOM	10112	CB	ILE A 122	-69.382	100.330	-32.336	1.00	31.44	T9
	ATOM	10113	CG2	ILE A 122	-69.964	99.035	-31.795	1.00	30.78	T9
	ATOM	10114	CG1	ILE A 122	-68.130	100.721	-31.548	1.00	24.52	T9
35	ATOM	10115	CD1	ILE A 122	-68.364	100.916	-30.077	1.00	33.98	T9
	ATOM	10116	C	ILE A 122	-70.307	99.843	-34.596	1.00	35.10	T9
	ATOM	10117	O	ILE A 122	-71.247	100.639	-34.612	1.00	38.05	T9
	ATOM	10118	N	PRO A 123	-70.351	98.682	-35.264	1.00	32.58	T9
	ATOM	10119	CD	PRO A 123	-69.228	97.743	-35.391	1.00	25.21	T9
40	ATOM	10120	CA	PRO A 123	-71.491	98.229	-36.070	1.00	27.35	T9
	ATOM	10121	CB	PRO A 123	-70.923	97.039	-36.845	1.00	29.12	T9
	ATOM	10122	CG	PRO A 123	-69.431	97.234	-36.782	1.00	31.06	T9
	ATOM	10123	C	PRO A 123	-72.710	97.819	-35.246	1.00	32.64	T9
	ATOM	10124	O	PRO A 123	-73.202	96.699	-35.375	1.00	28.99	T9
45	ATOM	10125	N	ARG A 124	-73.187	98.715	-34.392	1.00	20.82	T9
	ATOM	10126	CA	ARG A 124	-74.362	98.436	-33.577	1.00	29.65	T9
	ATOM	10127	CB	ARG A 124	-73.983	97.836	-32.246	1.00	25.26	T9
	ATOM	10128	CG	ARG A 124	-73.832	96.348	-32.297	1.00	31.10	T9
	ATOM	10129	CD	ARG A 124	-74.486	95.746	-31.068	1.00	33.12	T9
50	ATOM	10130	NE	ARG A 124	-75.945	95.795	-31.145	1.00	34.61	T9
	ATOM	10131	CZ	ARG A 124	-76.746	95.492	-30.126	1.00	35.58	T9
	ATOM	10132	NH1	ARG A 124	-76.217	95.131	-28.958	1.00	29.82	T9
	ATOM	10133	NH2	ARG A 124	-78.070	95.527	-30.276	1.00	29.61	T9
	ATOM	10134	C	ARG A 124	-75.100	99.717	-33.337	1.00	26.74	T9
55	ATOM	10135	O	ARG A 124	-74.506	100.795	-33.413	1.00	34.34	T9
	ATOM	10136	N	GLU A 125	-76.389	99.609	-33.029	1.00	31.24	T9
	ATOM	10137	CA	GLU A 125	-77.179	100.804	-32.823	1.00	34.85	T9
	ATOM	10138	CB	GLU A 125	-78.660	100.475	-32.947	1.00	33.64	T9
	ATOM	10139	CG	GLU A 125	-79.067	100.207	-34.394	1.00	34.13	T9
60	ATOM	10140	CD	GLU A 125	-80.412	100.843	-34.753	1.00	29.51	T9
	ATOM	10141	OE1	GLU A 125	-81.416	100.562	-34.038	1.00	25.06	T9
	ATOM	10142	OE2	GLU A 125	-80.455	101.620	-35.752	1.00	19.82	T9
	ATOM	10143	C	GLU A 125	-76.895	101.564	-31.537	1.00	29.00	T9
	ATOM	10144	O	GLU A 125	-76.686	102.781	-31.567	1.00	31.74	T9
65	ATOM	10145	N	ASN A 126	-76.895	100.881	-30.407	1.00	29.19	T9
	ATOM	10146	CA	ASN A 126	-76.592	101.582	-29.174	1.00	32.55	T9

	ATOM	10147	CB	ASN	A	126	-77.861	101.910	-28.380	1.00	26.22	T9
	ATOM	10148	CG	ASN	A	126	-78.554	103.165	-28.895	1.00	28.28	T9
	ATOM	10149	OD1	ASN	A	126	-79.266	103.129	-29.906	1.00	29.93	T9
	ATOM	10150	ND2	ASN	A	126	-78.328	104.295	-28.211	1.00	29.37	T9
5	ATOM	10151	C	ASN	A	126	-75.670	100.724	-28.363	1.00	34.78	T9
	ATOM	10152	O	ASN	A	126	-76.010	100.271	-27.271	1.00	33.68	T9
	ATOM	10153	N	ALA	A	127	-74.489	100.504	-28.922	1.00	32.42	T9
	ATOM	10154	CA	ALA	A	127	-73.484	99.683	-28.286	1.00	31.26	T9
	ATOM	10155	CB	ALA	A	127	-72.175	99.813	-29.047	1.00	30.99	T9
10	ATOM	10156	C	ALA	A	127	-73.282	100.043	-26.820	1.00	28.41	T9
	ATOM	10157	O	ALA	A	127	-73.163	101.214	-26.474	1.00	29.65	T9
	ATOM	10158	N	GLN	A	128	-73.271	99.027	-25.962	1.00	35.51	T9
	ATOM	10159	CA	GLN	A	128	-73.037	99.238	-24.541	1.00	37.43	T9
	ATOM	10160	CB	GLN	A	128	-73.663	98.115	-23.719	1.00	39.12	T9
15	ATOM	10161	CG	GLN	A	128	-75.179	98.118	-23.781	1.00	28.29	T9
	ATOM	10162	CD	GLN	A	128	-75.751	99.509	-23.578	1.00	30.95	T9
	ATOM	10163	OE1	GLN	A	128	-75.522	100.151	-22.545	1.00	32.04	T9
	ATOM	10164	NE2	GLN	A	128	-76.493	99.990	-24.572	1.00	25.85	T9
	ATOM	10165	C	GLN	A	128	-71.524	99.248	-24.384	1.00	26.94	T9
20	ATOM	10166	O	GLN	A	128	-70.876	98.200	-24.328	1.00	36.18	T9
	ATOM	10167	N	ILE	A	129	-70.979	100.457	-24.302	1.00	31.23	T9
	ATOM	10168	CA	ILE	A	129	-69.542	100.683	-24.232	1.00	30.50	T9
	ATOM	10169	CB	ILE	A	129	-69.136	101.483	-25.523	1.00	25.94	T9
	ATOM	10170	CG2	ILE	A	129	-68.317	102.703	-25.210	1.00	25.83	T9
25	ATOM	10171	CG1	ILE	A	129	-68.444	100.551	-26.492	1.00	28.33	T9
	ATOM	10172	CD1	ILE	A	129	-69.322	99.404	-26.911	1.00	30.67	T9
	ATOM	10173	C	ILE	A	129	-69.081	101.411	-22.968	1.00	25.29	T9
	ATOM	10174	O	ILE	A	129	-69.877	102.070	-22.302	1.00	32.16	T9
	ATOM	10175	N	SER	A	130	-67.806	101.267	-22.627	1.00	34.60	T9
30	ATOM	10176	CA	SER	A	130	-67.261	101.975	-21.483	1.00	33.30	T9
	ATOM	10177	CB	SER	A	130	-66.313	101.089	-20.690	1.00	26.24	T9
	ATOM	10178	OG	SER	A	130	-65.602	101.869	-19.737	1.00	26.50	T9
	ATOM	10179	C	SER	A	130	-66.485	103.166	-22.027	1.00	27.98	T9
	ATOM	10180	O	SER	A	130	-65.608	102.994	-22.870	1.00	33.54	T9
35	ATOM	10181	N	LEU	A	131	-66.793	104.372	-21.558	1.00	31.15	T9
	ATOM	10182	CA	LEU	A	131	-66.090	105.557	-22.046	1.00	29.74	T9
	ATOM	10183	CB	LEU	A	131	-67.070	106.707	-22.256	1.00	34.78	T9
	ATOM	10184	CG	LEU	A	131	-67.900	106.643	-23.531	1.00	31.89	T9
	ATOM	10185	CD1	LEU	A	131	-68.718	105.392	-23.554	1.00	37.60	T9
40	ATOM	10186	CD2	LEU	A	131	-68.794	107.846	-23.591	1.00	28.07	T9
	ATOM	10187	C	LEU	A	131	-64.936	106.032	-21.167	1.00	28.42	T9
	ATOM	10188	O	LEU	A	131	-64.662	107.227	-21.090	1.00	27.77	T9
	ATOM	10189	N	ASP	A	132	-64.256	105.100	-20.509	1.00	31.88	T9
	ATOM	10190	CA	ASP	A	132	-63.122	105.446	-19.660	1.00	35.42	T9
45	ATOM	10191	CB	ASP	A	132	-62.950	104.410	-18.547	1.00	35.35	T9
	ATOM	10192	CG	ASP	A	132	-63.872	104.660	-17.371	1.00	37.04	T9
	ATOM	10193	OD1	ASP	A	132	-63.980	103.776	-16.494	1.00	29.28	T9
	ATOM	10194	OD2	ASP	A	132	-64.480	105.751	-17.320	1.00	26.83	T9
	ATOM	10195	C	ASP	A	132	-61.822	105.569	-20.458	1.00	34.45	T9
50	ATOM	10196	O	ASP	A	132	-61.554	104.801	-21.384	1.00	25.41	T9
	ATOM	10197	N	GLY	A	133	-61.016	106.551	-20.082	1.00	33.30	T9
	ATOM	10198	CA	GLY	A	133	-59.753	106.786	-20.752	1.00	33.52	T9
	ATOM	10199	C	GLY	A	133	-58.829	105.589	-20.845	1.00	26.87	T9
	ATOM	10200	O	GLY	A	133	-58.079	105.500	-21.810	1.00	40.10	T9
55	ATOM	10201	N	ASP	A	134	-58.866	104.679	-19.870	1.00	31.93	T9
	ATOM	10202	CA	ASP	A	134	-57.992	103.511	-19.909	1.00	26.44	T9
	ATOM	10203	CB	ASP	A	134	-58.028	102.690	-18.626	1.00	33.05	T9
	ATOM	10204	CG	ASP	A	134	-58.423	103.468	-17.453	1.00	37.12	T9
	ATOM	10205	OD1	ASP	A	134	-57.662	104.381	-17.118	1.00	28.65	T9
60	ATOM	10206	OD2	ASP	A	134	-59.481	103.164	-16.865	1.00	35.09	T9
	ATOM	10207	C	ASP	A	134	-58.431	102.542	-20.959	1.00	32.30	T9
	ATOM	10208	O	ASP	A	134	-57.678	102.175	-21.832	1.00	31.71	T9
	ATOM	10209	N	VAL	A	135	-59.669	102.105	-20.829	1.00	27.74	T9
	ATOM	10210	CA	VAL	A	135	-60.220	101.096	-21.690	1.00	33.74	T9
65	ATOM	10211	CB	VAL	A	135	-61.477	100.547	-21.050	1.00	23.83	T9
	ATOM	10212	CG1	VAL	A	135	-61.122	99.930	-19.703	1.00	31.82	T9

	ATOM	10213	CG2	VAL	A	135	-62.491	101.656	-20.878	1.00	33.80	T9
	ATOM	10214	C	VAL	A	135	-60.474	101.374	-23.156	1.00	30.58	T9
	ATOM	10215	O	VAL	A	135	-60.269	100.486	-23.970	1.00	30.19	T9
	ATOM	10216	N	THR	A	136	-60.915	102.568	-23.523	1.00	34.53	T9
5	ATOM	10217	CA	THR	A	136	-61.144	102.806	-24.947	1.00	31.42	T9
	ATOM	10218	CB	THR	A	136	-62.670	102.878	-25.276	1.00	29.25	T9
	ATOM	10219	OG1	THR	A	136	-63.190	104.147	-24.894	1.00	27.49	T9
	ATOM	10220	CG2	THR	A	136	-63.430	101.814	-24.513	1.00	37.90	T9
	ATOM	10221	C	THR	A	136	-60.412	104.043	-25.497	1.00	35.94	T9
10	ATOM	10222	O	THR	A	136	-60.618	105.165	-25.039	1.00	33.33	T9
	ATOM	10223	N	PHE	A	137	-59.553	103.809	-26.487	1.00	34.65	T9
	ATOM	10224	CA	PHE	A	137	-58.752	104.860	-27.097	1.00	29.82	T9
	ATOM	10225	CB	PHE	A	137	-57.426	104.970	-26.347	1.00	28.16	T9
	ATOM	10226	CG	PHE	A	137	-56.788	103.643	-26.038	1.00	34.03	T9
15	ATOM	10227	CD1	PHE	A	137	-56.049	102.971	-26.996	1.00	29.99	T9
	ATOM	10228	CD2	PHE	A	137	-56.920	103.073	-24.785	1.00	34.64	T9
	ATOM	10229	CE1	PHE	A	137	-55.452	101.756	-26.706	1.00	32.10	T9
	ATOM	10230	CE2	PHE	A	137	-56.328	101.859	-24.488	1.00	28.93	T9
	ATOM	10231	CZ	PHE	A	137	-55.595	101.203	-25.445	1.00	26.63	T9
20	ATOM	10232	C	PHE	A	137	-58.507	104.636	-28.589	1.00	32.35	T9
	ATOM	10233	O	PHE	A	137	-58.702	103.539	-29.090	1.00	34.33	T9
	ATOM	10234	N	PHE	A	138	-58.062	105.675	-29.291	1.00	26.69	T9
	ATOM	10235	CA	PHE	A	138	-57.843	105.594	-30.729	1.00	25.68	T9
	ATOM	10236	CB	PHE	A	138	-58.948	106.402	-31.420	1.00	32.05	T9
25	ATOM	10237	CG	PHE	A	138	-59.069	106.157	-32.887	1.00	32.91	T9
	ATOM	10238	CD1	PHE	A	138	-58.604	104.983	-33.457	1.00	25.72	T9
	ATOM	10239	CD2	PHE	A	138	-59.649	107.113	-33.704	1.00	35.49	T9
	ATOM	10240	CE1	PHE	A	138	-58.710	104.767	-34.821	1.00	25.59	T9
	ATOM	10241	CE2	PHE	A	138	-59.760	106.909	-35.065	1.00	27.34	T9
30	ATOM	10242	CZ	PHE	A	138	-59.288	105.734	-35.626	1.00	30.68	T9
	ATOM	10243	C	PHE	A	138	-56.433	106.052	-31.138	1.00	34.92	T9
	ATOM	10244	O	PHE	A	138	-55.968	107.118	-30.751	1.00	35.88	T9
	ATOM	10245	N	GLY	A	139	-55.788	105.216	-31.955	1.00	29.59	T9
	ATOM	10246	CA	GLY	A	139	-54.407	105.390	-32.396	1.00	39.39	T9
35	ATOM	10247	C	GLY	A	139	-53.871	106.404	-33.378	1.00	31.92	T9
	ATOM	10248	O	GLY	A	139	-54.183	107.576	-33.275	1.00	30.03	T9
	ATOM	10249	N	ALA	A	140	-53.009	105.938	-34.287	1.00	29.45	T9
	ATOM	10250	CA	ALA	A	140	-52.350	106.749	-35.341	1.00	27.49	T9
	ATOM	10251	CB	ALA	A	140	-53.285	107.827	-35.847	1.00	32.20	T9
40	ATOM	10252	C	ALA	A	140	-50.993	107.381	-35.003	1.00	29.88	T9
	ATOM	10253	O	ALA	A	140	-50.914	108.368	-34.292	1.00	25.42	T9
	ATOM	10254	N	LEU	A	141	-49.933	106.796	-35.551	1.00	28.70	T9
	ATOM	10255	CA	LEU	A	141	-48.549	107.251	-35.366	1.00	28.58	T9
	ATOM	10256	CB	LEU	A	141	-47.838	106.382	-34.324	1.00	29.33	T9
45	ATOM	10257	CG	LEU	A	141	-46.334	106.573	-34.098	1.00	36.66	T9
	ATOM	10258	CD1	LEU	A	141	-45.926	105.978	-32.778	1.00	26.34	T9
	ATOM	10259	CD2	LEU	A	141	-45.563	105.914	-35.204	1.00	33.52	T9
	ATOM	10260	C	LEU	A	141	-47.823	107.136	-36.713	1.00	26.54	T9
	ATOM	10261	O	LEU	A	141	-47.925	106.116	-37.391	1.00	30.57	T9
50	ATOM	10262	N	LYS	A	142	-47.081	108.166	-37.101	1.00	31.75	T9
	ATOM	10263	CA	LYS	A	142	-46.387	108.128	-38.381	1.00	34.88	T9
	ATOM	10264	CB	LYS	A	142	-46.272	109.538	-38.949	1.00	21.76	T9
	ATOM	10265	CG	LYS	A	142	-45.562	109.562	-40.280	1.00	23.37	T9
	ATOM	10266	CD	LYS	A	142	-45.711	110.889	-40.993	1.00	27.64	T9
55	ATOM	10267	CE	LYS	A	142	-44.997	110.836	-42.338	1.00	34.42	T9
	ATOM	10268	NZ	LYS	A	142	-45.085	112.130	-43.069	1.00	23.94	T9
	ATOM	10269	C	LYS	A	142	-45.003	107.475	-38.371	1.00	25.80	T9
	ATOM	10270	O	LYS	A	142	-44.148	107.824	-37.562	1.00	30.34	T9
	ATOM	10271	N	LEU	A	143	-44.788	106.532	-39.284	1.00	28.94	T9
60	ATOM	10272	CA	LEU	A	143	-43.505	105.839	-39.391	1.00	30.98	T9
	ATOM	10273	CB	LEU	A	143	-43.684	104.498	-40.100	1.00	30.52	T9
	ATOM	10274	CG	LEU	A	143	-44.665	103.487	-39.503	1.00	29.24	T9
	ATOM	10275	CD1	LEU	A	143	-44.757	102.271	-40.409	1.00	39.58	T9
	ATOM	10276	CD2	LEU	A	143	-44.199	103.077	-38.114	1.00	26.58	T9
65	ATOM	10277	C	LEU	A	143	-42.510	106.682	-40.182	1.00	31.62	T9
	ATOM	10278	O	LEU	A	143	-42.902	107.529	-40.985	1.00	26.00	T9

	ATOM	10279	N	LEU	A	144	-41.222	106.452	-39.959	1.00	29.70	T9
	ATOM	10280	CA	LEU	A	144	-40.201	107.201	-40.681	1.00	37.07	T9
	ATOM	10281	CB	LEU	A	144	-38.928	107.326	-39.849	1.00	21.70	T9
	ATOM	10282	CG	LEU	A	144	-39.049	108.154	-38.569	1.00	23.26	T9
5	ATOM	10283	CD1	LEU	A	144	-37.768	108.024	-37.759	1.00	30.36	T9
	ATOM	10284	CD2	LEU	A	144	-39.321	109.615	-38.917	1.00	35.07	T9
	ATOM	10285	C	LEU	A	144	-39.870	106.500	-41.985	1.00	30.67	T9
	ATOM	10286	O	LEU	A	144	-40.210	105.307	-42.116	1.00	37.51	T9
	ATOM	10287	OXT	LEU	A	144	-39.257	107.154	-42.851	1.00	31.73	T9
10	ATOM	10288	CB	VAL	A	1	-4.896	131.372	-9.330	1.00	37.80	T10
	ATOM	10289	CG1	VAL	A	1	-5.845	132.571	-9.523	1.00	26.64	T10
	ATOM	10290	CG2	VAL	A	1	-4.397	130.875	-10.690	1.00	33.45	T10
	ATOM	10291	C	VAL	A	1	-4.205	132.152	-7.020	1.00	33.21	T10
	ATOM	10292	O	VAL	A	1	-4.029	133.293	-6.586	1.00	33.20	T10
15	ATOM	10293	N	VAL	A	1	-2.660	130.686	-8.358	1.00	38.53	T10
	ATOM	10294	CA	VAL	A	1	-3.679	131.777	-8.420	1.00	33.86	T10
	ATOM	10295	N	THR	A	2	-4.845	131.211	-6.317	1.00	31.67	T10
	ATOM	10296	CA	THR	A	2	-5.377	131.511	-4.984	1.00	27.68	T10
	ATOM	10297	CB	THR	A	2	-6.862	131.174	-4.877	1.00	34.41	T10
20	ATOM	10298	OG1	THR	A	2	-7.018	129.755	-4.878	1.00	26.81	T10
	ATOM	10299	CG2	THR	A	2	-7.630	131.767	-6.042	1.00	32.82	T10
	ATOM	10300	C	THR	A	2	-4.660	130.757	-3.879	1.00	32.99	T10
	ATOM	10301	O	THR	A	2	-3.845	129.881	-4.152	1.00	23.76	T10
	ATOM	10302	N	GLN	A	3	-4.978	131.094	-2.631	1.00	32.90	T10
25	ATOM	10303	CA	GLN	A	3	-4.364	130.444	-1.474	1.00	34.90	T10
	ATOM	10304	CB	GLN	A	3	-3.777	131.491	-0.530	1.00	34.63	T10
	ATOM	10305	CG	GLN	A	3	-2.968	132.555	-1.218	1.00	24.13	T10
	ATOM	10306	CD	GLN	A	3	-2.286	133.472	-0.226	1.00	27.36	T10
	ATOM	10307	OE1	GLN	A	3	-1.475	133.019	0.582	1.00	32.83	T10
30	ATOM	10308	NE2	GLN	A	3	-2.614	134.769	-0.276	1.00	30.22	T10
	ATOM	10309	C	GLN	A	3	-5.368	129.602	-0.692	1.00	34.95	T10
	ATOM	10310	O	GLN	A	3	-6.198	130.143	0.040	1.00	32.94	T10
	ATOM	10311	N	ASP	A	4	-5.294	128.284	-0.828	1.00	33.13	T10
	ATOM	10312	CA	ASP	A	4	-6.216	127.430	-0.093	1.00	30.33	T10
35	ATOM	10313	CB	ASP	A	4	-5.995	125.964	-0.454	1.00	33.88	T10
	ATOM	10314	CG	ASP	A	4	-6.282	125.679	-1.911	1.00	29.34	T10
	ATOM	10315	OD1	ASP	A	4	-6.957	126.517	-2.544	1.00	32.75	T10
	ATOM	10316	OD2	ASP	A	4	-5.851	124.620	-2.426	1.00	27.34	T10
	ATOM	10317	C	ASP	A	4	-6.025	127.613	1.403	1.00	34.90	T10
40	ATOM	10318	O	ASP	A	4	-4.918	127.879	1.865	1.00	28.97	T10
	ATOM	10319	N	CYS	A	5	-7.111	127.480	2.155	1.00	33.80	T10
	ATOM	10320	CA	CYS	A	5	-7.069	127.613	3.607	1.00	30.23	T10
	ATOM	10321	CB	CYS	A	5	-7.086	129.097	4.037	1.00	32.07	T10
	ATOM	10322	SG	CYS	A	5	-7.952	130.279	2.942	1.00	31.65	T10
45	ATOM	10323	C	CYS	A	5	-8.239	126.880	4.225	1.00	33.81	T10
	ATOM	10324	O	CYS	A	5	-9.292	126.793	3.627	1.00	28.02	T10
	ATOM	10325	N	LEU	A	6	-8.044	126.324	5.410	1.00	30.02	T10
	ATOM	10326	CA	LEU	A	6	-9.110	125.609	6.106	1.00	31.74	T10
	ATOM	10327	CB	LEU	A	6	-8.941	124.097	5.946	1.00	25.93	T10
50	ATOM	10328	CG	LEU	A	6	-9.954	123.214	6.670	1.00	29.37	T10
	ATOM	10329	CD1	LEU	A	6	-10.148	121.931	5.908	1.00	31.78	T10
	ATOM	10330	CD2	LEU	A	6	-9.481	122.936	8.082	1.00	31.43	T10
	ATOM	10331	C	LEU	A	6	-9.028	126.001	7.570	1.00	32.29	T10
	ATOM	10332	O	LEU	A	6	-7.944	126.017	8.141	1.00	34.64	T10
55	ATOM	10333	N	GLN	A	7	-10.166	126.328	8.174	1.00	33.83	T10
	ATOM	10334	CA	GLN	A	7	-10.176	126.742	9.564	1.00	30.42	T10
	ATOM	10335	CB	GLN	A	7	-10.454	128.232	9.650	1.00	32.65	T10
	ATOM	10336	CG	GLN	A	7	-10.230	128.825	11.020	1.00	39.76	T10
	ATOM	10337	CD	GLN	A	7	-10.304	130.331	10.992	1.00	25.84	T10
60	ATOM	10338	OE1	GLN	A	7	-11.371	130.905	10.794	1.00	24.73	T10
	ATOM	10339	NE2	GLN	A	7	-9.165	130.982	11.167	1.00	34.01	T10
	ATOM	10340	C	GLN	A	7	-11.200	125.983	10.381	1.00	27.74	T10
	ATOM	10341	O	GLN	A	7	-12.300	125.711	9.912	1.00	32.82	T10
	ATOM	10342	N	LEU	A	8	-10.821	125.649	11.611	1.00	26.72	T10
65	ATOM	10343	CA	LEU	A	8	-11.686	124.913	12.517	1.00	28.61	T10
	ATOM	10344	CB	LEU	A	8	-11.009	123.608	12.953	1.00	26.29	T10

	ATOM	10345	CG	LEU	A	8	-11.128	122.342	12.104	1.00	32.96	T10
	ATOM	10346	CD1	LEU	A	8	-11.640	122.662	10.724	1.00	29.86	T10
	ATOM	10347	CD2	LEU	A	8	-9.775	121.672	12.039	1.00	30.43	T10
	ATOM	10348	C	LEU	A	8	-12.045	125.741	13.741	1.00	38.00	T10
5	ATOM	10349	O	LEU	A	8	-11.310	126.631	14.145	1.00	33.52	T10
	ATOM	10350	N	ILE	A	9	-13.190	125.422	14.327	1.00	32.32	T10
	ATOM	10351	CA	ILE	A	9	-13.709	126.105	15.508	1.00	31.15	T10
	ATOM	10352	CB	ILE	A	9	-14.977	126.896	15.146	1.00	33.79	T10
	ATOM	10353	CG2	ILE	A	9	-15.769	127.220	16.389	1.00	22.71	T10
10	ATOM	10354	CG1	ILE	A	9	-14.624	128.181	14.425	1.00	29.17	T10
	ATOM	10355	CD1	ILE	A	9	-15.860	128.937	14.048	1.00	27.87	T10
	ATOM	10356	C	ILE	A	9	-14.102	125.077	16.568	1.00	27.53	T10
	ATOM	10357	O	ILE	A	9	-14.607	124.005	16.244	1.00	26.30	T10
	ATOM	10358	N	ALA	A	10	-13.901	125.406	17.835	1.00	34.67	T10
15	ATOM	10359	CA	ALA	A	10	-14.275	124.472	18.888	1.00	29.64	T10
	ATOM	10360	CB	ALA	A	10	-13.889	125.028	20.241	1.00	32.62	T10
	ATOM	10361	C	ALA	A	10	-15.776	124.184	18.858	1.00	36.42	T10
	ATOM	10362	O	ALA	A	10	-16.598	125.099	18.789	1.00	34.96	T10
	ATOM	10363	N	ASP	A	11	-16.128	122.905	18.909	1.00	31.97	T10
20	ATOM	10364	CA	ASP	A	11	-17.526	122.505	18.915	1.00	32.66	T10
	ATOM	10365	CB	ASP	A	11	-17.710	121.177	18.191	1.00	35.55	T10
	ATOM	10366	CG	ASP	A	11	-19.126	120.666	18.292	1.00	32.17	T10
	ATOM	10367	OD1	ASP	A	11	-20.054	121.510	18.233	1.00	36.35	T10
	ATOM	10368	OD2	ASP	A	11	-19.317	119.434	18.423	1.00	25.15	T10
25	ATOM	10369	C	ASP	A	11	-18.015	122.384	20.355	1.00	23.21	T10
	ATOM	10370	O	ASP	A	11	-17.909	121.327	20.980	1.00	30.23	T10
	ATOM	10371	N	SER	A	12	-18.544	123.496	20.860	1.00	29.92	T10
	ATOM	10372	CA	SER	A	12	-19.051	123.608	22.225	1.00	29.48	T10
	ATOM	10373	CB	SER	A	12	-19.454	125.058	22.520	1.00	32.24	T10
30	ATOM	10374	OG	SER	A	12	-20.452	125.528	21.615	1.00	33.45	T10
	ATOM	10375	C	SER	A	12	-20.234	122.712	22.508	1.00	29.30	T10
	ATOM	10376	O	SER	A	12	-20.994	122.981	23.423	1.00	31.69	T10
	ATOM	10377	N	GLU	A	13	-20.402	121.652	21.728	1.00	31.63	T10
	ATOM	10378	CA	GLU	A	13	-21.527	120.760	21.960	1.00	32.54	T10
35	ATOM	10379	CB	GLU	A	13	-22.654	121.079	20.999	1.00	27.00	T10
	ATOM	10380	CG	GLU	A	13	-23.594	122.102	21.578	1.00	36.74	T10
	ATOM	10381	CD	GLU	A	13	-24.764	122.365	20.662	1.00	25.29	T10
	ATOM	10382	OE1	GLU	A	13	-25.300	121.370	20.107	1.00	37.44	T10
	ATOM	10383	OE2	GLU	A	13	-25.158	123.552	20.491	1.00	34.66	T10
40	ATOM	10384	C	GLU	A	13	-21.233	119.279	21.925	1.00	29.37	T10
	ATOM	10385	O	GLU	A	13	-22.118	118.468	21.635	1.00	31.62	T10
	ATOM	10386	N	THR	A	14	-19.982	118.935	22.208	1.00	32.05	T10
	ATOM	10387	CA	THR	A	14	-19.557	117.548	22.269	1.00	30.26	T10
	ATOM	10388	CB	THR	A	14	-19.038	116.996	20.920	1.00	33.26	T10
45	ATOM	10389	OG1	THR	A	14	-17.952	117.799	20.465	1.00	28.60	T10
	ATOM	10390	CG2	THR	A	14	-20.153	116.985	19.872	1.00	27.19	T10
	ATOM	10391	C	THR	A	14	-18.447	117.531	23.290	1.00	32.21	T10
	ATOM	10392	O	THR	A	14	-17.744	118.521	23.481	1.00	31.91	T10
	ATOM	10393	N	PRO	A	15	-18.289	116.403	23.981	1.00	27.59	T10
50	ATOM	10394	CD	PRO	A	15	-19.057	115.159	23.813	1.00	34.37	T10
	ATOM	10395	CA	PRO	A	15	-17.259	116.247	25.011	1.00	36.06	T10
	ATOM	10396	CB	PRO	A	15	-17.511	114.831	25.547	1.00	20.86	T10
	ATOM	10397	CG	PRO	A	15	-18.954	114.559	25.186	1.00	27.27	T10
	ATOM	10398	C	PRO	A	15	-15.865	116.376	24.418	1.00	28.56	T10
55	ATOM	10399	O	PRO	A	15	-15.631	115.970	23.277	1.00	28.37	T10
	ATOM	10400	N	THR	A	16	-14.941	116.932	25.190	1.00	27.05	T10
	ATOM	10401	CA	THR	A	16	-13.577	117.073	24.715	1.00	27.07	T10
	ATOM	10402	CB	THR	A	16	-12.790	117.968	25.653	1.00	34.68	T10
	ATOM	10403	OG1	THR	A	16	-12.580	117.297	26.905	1.00	24.83	T10
60	ATOM	10404	CG2	THR	A	16	-13.577	119.237	25.912	1.00	30.13	T10
	ATOM	10405	C	THR	A	16	-12.942	115.690	24.690	1.00	34.26	T10
	ATOM	10406	O	THR	A	16	-12.791	115.057	25.730	1.00	27.88	T10
	ATOM	10407	N	ILE	A	17	-12.573	115.222	23.506	1.00	31.76	T10
	ATOM	10408	CA	ILE	A	17	-11.966	113.897	23.355	1.00	30.44	T10
65	ATOM	10409	CB	ILE	A	17	-11.510	113.676	21.908	1.00	27.51	T10
	ATOM	10410	CG2	ILE	A	17	-10.867	112.320	21.772	1.00	27.37	T10

	ATOM	10411	CG1	ILE	A	17	-12.704	113.798	20.968	1.00	32.31	T10
	ATOM	10412	CD1	ILE	A	17	-12.353	113.652	19.525	1.00	29.91	T10
	ATOM	10413	C	ILE	A	17	-10.772	113.608	24.278	1.00	26.91	T10
	ATOM	10414	O	ILE	A	17	-9.837	114.410	24.390	1.00	35.92	T10
5	ATOM	10415	N	GLN	A	18	-10.818	112.450	24.935	1.00	32.62	T10
	ATOM	10416	CA	GLN	A	18	-9.750	112.022	25.843	1.00	24.64	T10
	ATOM	10417	CB	GLN	A	18	-10.308	111.694	27.218	1.00	33.46	T10
	ATOM	10418	CG	GLN	A	18	-9.643	112.480	28.303	1.00	31.56	T10
	ATOM	10419	CD	GLN	A	18	-10.086	113.924	28.291	1.00	30.27	T10
10	ATOM	10420	OE1	GLN	A	18	-11.194	114.248	28.720	1.00	33.16	T10
	ATOM	10421	NE2	GLN	A	18	-9.230	114.803	27.785	1.00	28.12	T10
	ATOM	10422	C	GLN	A	18	-9.025	110.790	25.302	1.00	29.84	T10
	ATOM	10423	O	GLN	A	18	-9.656	109.824	24.860	1.00	25.41	T10
	ATOM	10424	N	LYS	A	19	-7.698	110.811	25.351	1.00	28.29	T10
15	ATOM	10425	CA	LYS	A	19	-6.922	109.693	24.832	1.00	32.32	T10
	ATOM	10426	CB	LYS	A	19	-7.127	109.585	23.319	1.00	22.96	T10
	ATOM	10427	CG	LYS	A	19	-6.167	108.636	22.626	1.00	30.43	T10
	ATOM	10428	CD	LYS	A	19	-6.565	108.424	21.166	1.00	31.18	T10
	ATOM	10429	CE	LYS	A	19	-5.614	107.459	20.454	1.00	24.21	T10
20	ATOM	10430	NZ	LYS	A	19	-6.016	107.160	19.041	1.00	35.27	T10
	ATOM	10431	C	LYS	A	19	-5.435	109.818	25.138	1.00	31.64	T10
	ATOM	10432	O	LYS	A	19	-4.824	110.864	24.897	1.00	30.29	T10
	ATOM	10433	N	GLY	A	20	-4.858	108.734	25.653	1.00	24.10	T10
	ATOM	10434	CA	GLY	A	20	-3.447	108.730	25.995	1.00	36.72	T10
25	ATOM	10435	C	GLY	A	20	-3.132	109.880	26.930	1.00	31.13	T10
	ATOM	10436	O	GLY	A	20	-2.091	110.528	26.786	1.00	30.28	T10
	ATOM	10437	N	SER	A	21	-4.038	110.132	27.880	1.00	29.03	T10
	ATOM	10438	CA	SER	A	21	-3.888	111.222	28.853	1.00	29.92	T10
	ATOM	10439	CB	SER	A	21	-2.789	110.886	29.880	1.00	29.51	T10
30	ATOM	10440	OG	SER	A	21	-1.525	110.701	29.260	1.00	27.89	T10
	ATOM	10441	C	SER	A	21	-3.597	112.576	28.182	1.00	30.13	T10
	ATOM	10442	O	SER	A	21	-2.787	113.384	28.665	1.00	27.89	T10
	ATOM	10443	N	TYR	A	22	-4.272	112.792	27.055	1.00	25.15	T10
	ATOM	10444	CA	TYR	A	22	-4.165	114.019	26.268	1.00	32.81	T10
35	ATOM	10445	CB	TYR	A	22	-3.511	113.734	24.918	1.00	27.86	T10
	ATOM	10446	CG	TYR	A	22	-2.032	113.993	24.872	1.00	29.05	T10
	ATOM	10447	CD1	TYR	A	22	-1.261	113.949	26.032	1.00	25.11	T10
	ATOM	10448	CE1	TYR	A	22	0.125	114.155	25.988	1.00	33.27	T10
	ATOM	10449	CD2	TYR	A	22	-1.390	114.247	23.663	1.00	36.52	T10
40	ATOM	10450	CE2	TYR	A	22	-0.009	114.448	23.606	1.00	28.83	T10
	ATOM	10451	CZ	TYR	A	22	0.745	114.401	24.772	1.00	32.67	T10
	ATOM	10452	OH	TYR	A	22	2.113	114.591	24.725	1.00	26.58	T10
	ATOM	10453	C	TYR	A	22	-5.589	114.492	26.025	1.00	27.11	T10
	ATOM	10454	O	TYR	A	22	-6.504	113.669	25.897	1.00	35.66	T10
45	ATOM	10455	N	THR	A	23	-5.790	115.805	25.971	1.00	31.58	T10
	ATOM	10456	CA	THR	A	23	-7.128	116.318	25.716	1.00	30.59	T10
	ATOM	10457	CB	THR	A	23	-7.522	117.431	26.700	1.00	32.60	T10
	ATOM	10458	OG1	THR	A	23	-7.076	117.100	28.019	1.00	30.08	T10
	ATOM	10459	CG2	THR	A	23	-9.031	117.567	26.729	1.00	32.49	T10
50	ATOM	10460	C	THR	A	23	-7.164	116.874	24.303	1.00	30.30	T10
	ATOM	10461	O	THR	A	23	-6.285	117.637	23.906	1.00	26.08	T10
	ATOM	10462	N	PHE	A	24	-8.170	116.473	23.539	1.00	30.85	T10
	ATOM	10463	CA	PHE	A	24	-8.302	116.945	22.169	1.00	27.24	T10
	ATOM	10464	CB	PHE	A	24	-8.254	115.770	21.192	1.00	27.52	T10
55	ATOM	10465	CG	PHE	A	24	-6.930	115.068	21.154	1.00	27.36	T10
	ATOM	10466	CD1	PHE	A	24	-6.584	114.145	22.130	1.00	31.35	T10
	ATOM	10467	CD2	PHE	A	24	-6.018	115.346	20.150	1.00	24.80	T10
	ATOM	10468	CE1	PHE	A	24	-5.346	113.513	22.102	1.00	29.43	T10
	ATOM	10469	CE2	PHE	A	24	-4.781	114.721	20.117	1.00	35.64	T10
60	ATOM	10470	CZ	PHE	A	24	-4.446	113.804	21.094	1.00	30.82	T10
	ATOM	10471	C	PHE	A	24	-9.598	117.730	21.969	1.00	29.37	T10
	ATOM	10472	O	PHE	A	24	-10.693	117.240	22.259	1.00	24.85	T10
	ATOM	10473	N	VAL	A	25	-9.464	118.954	21.475	1.00	31.68	T10
	ATOM	10474	CA	VAL	A	25	-10.608	119.816	21.230	1.00	31.43	T10
65	ATOM	10475	CB	VAL	A	25	-10.153	121.217	20.796	1.00	28.55	T10
	ATOM	10476	CG1	VAL	A	25	-11.344	122.072	20.462	1.00	26.72	T10

	ATOM	10477	CG2	VAL	A	25	-9.337	121.859	21.891	1.00	22.72	T10
	ATOM	10478	C	VAL	A	25	-11.487	119.248	20.129	1.00	34.71	T10
	ATOM	10479	O	VAL	A	25	-10.998	118.815	19.092	1.00	26.23	T10
	ATOM	10480	N	PRO	A	26	-12.804	119.228	20.347	1.00	33.12	T10
5	ATOM	10481	CD	PRO	A	26	-13.466	119.513	21.625	1.00	29.27	T10
	ATOM	10482	CA	PRO	A	26	-13.762	118.713	19.361	1.00	25.71	T10
	ATOM	10483	CB	PRO	A	26	-15.041	118.549	20.168	1.00	32.08	T10
	ATOM	10484	CG	PRO	A	26	-14.572	118.515	21.605	1.00	31.68	T10
	ATOM	10485	C	PRO	A	26	-13.926	119.794	18.300	1.00	29.42	T10
10	ATOM	10486	O	PRO	A	26	-14.491	120.848	18.583	1.00	28.66	T10
	ATOM	10487	N	TRP	A	27	-13.447	119.554	17.086	1.00	28.46	T10
	ATOM	10488	CA	TRP	A	27	-13.551	120.571	16.048	1.00	36.79	T10
	ATOM	10489	CB	TRP	A	27	-12.381	120.450	15.075	1.00	28.64	T10
	ATOM	10490	CG	TRP	A	27	-11.071	120.695	15.717	1.00	26.26	T10
15	ATOM	10491	CD2	TRP	A	27	-10.690	121.848	16.480	1.00	31.36	T10
	ATOM	10492	CE2	TRP	A	27	-9.375	121.627	16.941	1.00	29.54	T10
	ATOM	10493	CE3	TRP	A	27	-11.333	123.044	16.822	1.00	27.64	T10
	ATOM	10494	CD1	TRP	A	27	-10.002	119.850	15.738	1.00	38.85	T10
	ATOM	10495	NE1	TRP	A	27	-8.977	120.400	16.473	1.00	26.71	T10
20	ATOM	10496	CZ2	TRP	A	27	-8.689	122.560	17.729	1.00	27.98	T10
	ATOM	10497	CZ3	TRP	A	27	-10.649	123.968	17.605	1.00	29.33	T10
	ATOM	10498	CH2	TRP	A	27	-9.342	123.719	18.049	1.00	28.27	T10
	ATOM	10499	C	TRP	A	27	-14.852	120.605	15.264	1.00	29.33	T10
	ATOM	10500	O	TRP	A	27	-15.647	119.665	15.273	1.00	29.53	T10
25	ATOM	10501	N	LEU	A	28	-15.046	121.724	14.579	1.00	35.47	T10
	ATOM	10502	CA	LEU	A	28	-16.218	121.965	13.753	1.00	35.58	T10
	ATOM	10503	CB	LEU	A	28	-17.307	122.636	14.579	1.00	33.17	T10
	ATOM	10504	CG	LEU	A	28	-18.712	122.526	13.995	1.00	33.79	T10
	ATOM	10505	CD1	LEU	A	28	-19.122	121.051	13.955	1.00	29.72	T10
30	ATOM	10506	CD2	LEU	A	28	-19.676	123.339	14.850	1.00	25.60	T10
	ATOM	10507	C	LEU	A	28	-15.755	122.901	12.641	1.00	30.38	T10
	ATOM	10508	O	LEU	A	28	-15.149	123.939	12.903	1.00	29.90	T10
	ATOM	10509	N	LEU	A	29	-16.035	122.540	11.399	1.00	29.08	T10
	ATOM	10510	CA	LEU	A	29	-15.592	123.356	10.286	1.00	38.47	T10
35	ATOM	10511	CB	LEU	A	29	-16.094	122.793	8.968	1.00	27.68	T10
	ATOM	10512	CG	LEU	A	29	-15.674	123.638	7.771	1.00	27.22	T10
	ATOM	10513	CD1	LEU	A	29	-14.219	123.383	7.469	1.00	31.81	T10
	ATOM	10514	CD2	LEU	A	29	-16.513	123.300	6.578	1.00	27.38	T10
	ATOM	10515	C	LEU	A	29	-16.044	124.788	10.384	1.00	28.08	T10
40	ATOM	10516	O	LEU	A	29	-17.233	125.061	10.538	1.00	33.23	T10
	ATOM	10517	N	SER	A	30	-15.087	125.706	10.301	1.00	32.53	T10
	ATOM	10518	CA	SER	A	30	-15.394	127.130	10.327	1.00	24.73	T10
	ATOM	10519	CB	SER	A	30	-14.220	127.934	10.863	1.00	30.14	T10
	ATOM	10520	OG	SER	A	30	-14.497	129.314	10.771	1.00	28.70	T10
45	ATOM	10521	C	SER	A	30	-15.628	127.480	8.874	1.00	34.77	T10
	ATOM	10522	O	SER	A	30	-16.664	128.014	8.516	1.00	28.27	T10
	ATOM	10523	N	PHE	A	31	-14.652	127.156	8.039	1.00	35.72	T10
	ATOM	10524	CA	PHE	A	31	-14.754	127.401	6.608	1.00	28.12	T10
	ATOM	10525	CB	PHE	A	31	-14.681	128.902	6.309	1.00	24.19	T10
50	ATOM	10526	CG	PHE	A	31	-13.284	129.430	6.160	1.00	21.44	T10
	ATOM	10527	CD1	PHE	A	31	-12.622	129.336	4.940	1.00	33.85	T10
	ATOM	10528	CD2	PHE	A	31	-12.623	130.011	7.242	1.00	29.08	T10
	ATOM	10529	CE1	PHE	A	31	-11.327	129.811	4.798	1.00	36.55	T10
	ATOM	10530	CE2	PHE	A	31	-11.331	130.487	7.110	1.00	27.54	T10
55	ATOM	10531	CZ	PHE	A	31	-10.679	130.387	5.885	1.00	28.77	T10
	ATOM	10532	C	PHE	A	31	-13.623	126.659	5.902	1.00	32.07	T10
	ATOM	10533	O	PHE	A	31	-12.580	126.393	6.493	1.00	26.99	T10
	ATOM	10534	N	LYS	A	32	-13.844	126.312	4.643	1.00	33.02	T10
	ATOM	10535	CA	LYS	A	32	-12.848	125.604	3.859	1.00	33.32	T10
60	ATOM	10536	CB	LYS	A	32	-13.224	124.139	3.724	1.00	29.00	T10
	ATOM	10537	CG	LYS	A	32	-12.422	123.432	2.678	1.00	24.19	T10
	ATOM	10538	CD	LYS	A	32	-12.931	122.038	2.406	1.00	31.92	T10
	ATOM	10539	CE	LYS	A	32	-12.231	121.499	1.175	1.00	28.85	T10
	ATOM	10540	NZ	LYS	A	32	-12.548	120.072	0.948	1.00	25.66	T10
65	ATOM	10541	C	LYS	A	32	-12.783	126.242	2.486	1.00	36.42	T10
	ATOM	10542	O	LYS	A	32	-13.793	126.370	1.810	1.00	24.44	T10

	ATOM	10543	N	ARG	A	33	-11.590	126.637	2.074	1.00	31.33	T10
	ATOM	10544	CA	ARG	A	33	-11.396	127.294	0.792	1.00	26.03	T10
	ATOM	10545	CB	ARG	A	33	-11.035	128.762	1.038	1.00	26.82	T10
	ATOM	10546	CG	ARG	A	33	-10.640	129.573	-0.173	1.00	31.34	T10
5	ATOM	10547	CD	ARG	A	33	-10.805	131.054	0.145	1.00	33.49	T10
	ATOM	10548	NE	ARG	A	33	-10.361	131.932	-0.935	1.00	29.79	T10
	ATOM	10549	CZ	ARG	A	33	-9.095	132.274	-1.134	1.00	35.55	T10
	ATOM	10550	NH1	ARG	A	33	-8.150	131.819	-0.322	1.00	34.41	T10
	ATOM	10551	NH2	ARG	A	33	-8.771	133.059	-2.146	1.00	34.90	T10
10	ATOM	10552	C	ARG	A	33	-10.304	126.593	0.000	1.00	29.75	T10
	ATOM	10553	O	ARG	A	33	-9.170	126.447	0.469	1.00	24.98	T10
	ATOM	10554	N	GLY	A	34	-10.651	126.146	-1.198	1.00	23.95	T10
	ATOM	10555	CA	GLY	A	34	-9.675	125.468	-2.021	1.00	26.97	T10
	ATOM	10556	C	GLY	A	34	-9.697	123.961	-1.871	1.00	27.51	T10
15	ATOM	10557	O	GLY	A	34	-10.635	123.384	-1.316	1.00	29.48	T10
	ATOM	10558	N	SER	A	35	-8.637	123.322	-2.349	1.00	28.28	T10
	ATOM	10559	CA	SER	A	35	-8.539	121.872	-2.306	1.00	32.77	T10
	ATOM	10560	CB	SER	A	35	-8.456	121.345	-3.734	1.00	30.97	T10
	ATOM	10561	OG	SER	A	35	-7.371	121.958	-4.418	1.00	33.25	T10
20	ATOM	10562	C	SER	A	35	-7.366	121.301	-1.509	1.00	22.68	T10
	ATOM	10563	O	SER	A	35	-7.411	120.141	-1.110	1.00	29.10	T10
	ATOM	10564	N	ALA	A	36	-6.328	122.095	-1.279	1.00	33.02	T10
	ATOM	10565	CA	ALA	A	36	-5.149	121.608	-0.577	1.00	31.96	T10
	ATOM	10566	CB	ALA	A	36	-4.067	122.665	-0.605	1.00	30.46	T10
25	ATOM	10567	C	ALA	A	36	-5.346	121.119	0.848	1.00	26.81	T10
	ATOM	10568	O	ALA	A	36	-4.483	120.421	1.371	1.00	36.38	T10
	ATOM	10569	N	LEU	A	37	-6.463	121.469	1.483	1.00	26.01	T10
	ATOM	10570	CA	LEU	A	37	-6.710	121.046	2.862	1.00	32.36	T10
	ATOM	10571	CB	LEU	A	37	-6.420	122.210	3.807	1.00	34.45	T10
30	ATOM	10572	CG	LEU	A	37	-4.984	122.735	3.759	1.00	25.14	T10
	ATOM	10573	CD1	LEU	A	37	-4.901	124.122	4.350	1.00	30.96	T10
	ATOM	10574	CD2	LEU	A	37	-4.076	121.776	4.492	1.00	37.05	T10
	ATOM	10575	C	LEU	A	37	-8.126	120.520	3.092	1.00	37.28	T10
	ATOM	10576	O	LEU	A	37	-9.073	120.980	2.466	1.00	32.88	T10
35	ATOM	10577	N	GLU	A	38	-8.259	119.557	4.000	1.00	33.62	T10
	ATOM	10578	CA	GLU	A	38	-9.546	118.936	4.313	1.00	35.54	T10
	ATOM	10579	CB	GLU	A	38	-9.729	117.667	3.487	1.00	27.88	T10
	ATOM	10580	CG	GLU	A	38	-10.225	117.870	2.075	1.00	33.07	T10
	ATOM	10581	CD	GLU	A	38	-10.144	116.587	1.249	1.00	34.68	T10
40	ATOM	10582	OE1	GLU	A	38	-10.360	115.489	1.823	1.00	31.91	T10
	ATOM	10583	OE2	GLU	A	38	-9.873	116.685	0.026	1.00	34.18	T10
	ATOM	10584	C	GLU	A	38	-9.668	118.539	5.780	1.00	28.16	T10
	ATOM	10585	O	GLU	A	38	-8.672	118.466	6.492	1.00	30.02	T10
	ATOM	10586	N	GLU	A	39	-10.892	118.275	6.229	1.00	32.81	T10
45	ATOM	10587	CA	GLU	A	39	-11.105	117.841	7.602	1.00	31.23	T10
	ATOM	10588	CB	GLU	A	39	-12.479	118.209	8.125	1.00	26.59	T10
	ATOM	10589	CG	GLU	A	39	-12.953	119.578	7.822	1.00	30.35	T10
	ATOM	10590	CD	GLU	A	39	-14.468	119.606	7.743	1.00	31.88	T10
	ATOM	10591	OE1	GLU	A	39	-14.998	119.528	6.601	1.00	33.65	T10
50	ATOM	10592	OE2	GLU	A	39	-15.122	119.679	8.822	1.00	25.48	T10
	ATOM	10593	C	GLU	A	39	-11.092	116.337	7.539	1.00	32.56	T10
	ATOM	10594	O	GLU	A	39	-11.556	115.754	6.565	1.00	28.61	T10
	ATOM	10595	N	LYS	A	40	-10.588	115.702	8.579	1.00	37.41	T10
	ATOM	10596	CA	LYS	A	40	-10.582	114.259	8.596	1.00	32.73	T10
55	ATOM	10597	CB	LYS	A	40	-9.458	113.702	7.719	1.00	30.07	T10
	ATOM	10598	CG	LYS	A	40	-9.449	112.173	7.676	1.00	37.86	T10
	ATOM	10599	CD	LYS	A	40	-8.135	111.644	7.150	1.00	33.22	T10
	ATOM	10600	CE	LYS	A	40	-8.049	110.131	7.290	1.00	30.24	T10
	ATOM	10601	NZ	LYS	A	40	-6.679	109.641	6.914	1.00	32.58	T10
60	ATOM	10602	C	LYS	A	40	-10.429	113.757	10.019	1.00	32.30	T10
	ATOM	10603	O	LYS	A	40	-9.346	113.804	10.600	1.00	32.82	T10
	ATOM	10604	N	GLU	A	41	-11.534	113.294	10.582	1.00	32.50	T10
	ATOM	10605	CA	GLU	A	41	-11.520	112.768	11.929	1.00	27.31	T10
	ATOM	10606	CB	GLU	A	41	-10.830	111.409	11.915	1.00	34.58	T10
65	ATOM	10607	CG	GLU	A	41	-11.416	110.499	10.847	1.00	29.02	T10
	ATOM	10608	CD	GLU	A	41	-10.548	109.286	10.547	1.00	24.36	T10

	ATOM	10609	OE1	GLU	A	41	-9.331	109.465	10.278	1.00	26.08	T10
	ATOM	10610	OE2	GLU	A	41	-11.086	108.155	10.567	1.00	32.70	T10
	ATOM	10611	C	GLU	A	41	-10.834	113.713	12.903	1.00	28.08	T10
	ATOM	10612	O	GLU	A	41	-9.839	113.363	13.544	1.00	19.81	T10
5	ATOM	10613	N	ASN	A	42	-11.372	114.920	12.991	1.00	29.07	T10
	ATOM	10614	CA	ASN	A	42	-10.864	115.935	13.898	1.00	26.09	T10
	ATOM	10615	CB	ASN	A	42	-10.985	115.451	15.333	1.00	31.87	T10
	ATOM	10616	CG	ASN	A	42	-11.490	116.525	16.249	1.00	30.55	T10
	ATOM	10617	OD1	ASN	A	42	-10.876	116.821	17.267	1.00	33.34	T10
10	ATOM	10618	ND2	ASN	A	42	-12.622	117.124	15.892	1.00	30.37	T10
	ATOM	10619	C	ASN	A	42	-9.442	116.389	13.655	1.00	38.41	T10
	ATOM	10620	O	ASN	A	42	-8.813	116.937	14.553	1.00	28.43	T10
	ATOM	10621	N	LYS	A	43	-8.943	116.164	12.448	1.00	29.45	T10
	ATOM	10622	CA	LYS	A	43	-7.594	116.571	12.089	1.00	37.14	T10
15	ATOM	10623	CB	LYS	A	43	-6.668	115.359	12.028	1.00	32.83	T10
	ATOM	10624	CG	LYS	A	43	-6.372	114.740	13.375	1.00	29.86	T10
	ATOM	10625	CD	LYS	A	43	-5.509	113.501	13.240	1.00	28.86	T10
	ATOM	10626	CE	LYS	A	43	-6.294	112.346	12.615	1.00	27.96	T10
	ATOM	10627	NZ	LYS	A	43	-5.491	111.086	12.489	1.00	29.44	T10
20	ATOM	10628	C	LYS	A	43	-7.634	117.238	10.728	1.00	28.31	T10
	ATOM	10629	O	LYS	A	43	-8.597	117.067	9.974	1.00	20.67	T10
	ATOM	10630	N	ILE	A	44	-6.598	118.004	10.408	1.00	27.11	T10
	ATOM	10631	CA	ILE	A	44	-6.541	118.655	9.109	1.00	30.55	T10
	ATOM	10632	CB	ILE	A	44	-5.937	120.054	9.213	1.00	38.41	T10
25	ATOM	10633	CG2	ILE	A	44	-5.894	120.694	7.847	1.00	29.49	T10
	ATOM	10634	CG1	ILE	A	44	-6.775	120.907	10.161	1.00	27.20	T10
	ATOM	10635	CD1	ILE	A	44	-6.254	122.300	10.353	1.00	28.24	T10
	ATOM	10636	C	ILE	A	44	-5.678	117.799	8.194	1.00	32.72	T10
	ATOM	10637	O	ILE	A	44	-4.531	117.495	8.509	1.00	37.94	T10
30	ATOM	10638	N	LEU	A	45	-6.239	117.407	7.062	1.00	27.04	T10
	ATOM	10639	CA	LEU	A	45	-5.527	116.569	6.116	1.00	30.49	T10
	ATOM	10640	CB	LEU	A	45	-6.476	115.514	5.558	1.00	37.22	T10
	ATOM	10641	CG	LEU	A	45	-5.858	114.622	4.484	1.00	32.56	T10
	ATOM	10642	CD1	LEU	A	45	-4.826	113.691	5.115	1.00	31.89	T10
35	ATOM	10643	CD2	LEU	A	45	-6.950	113.836	3.811	1.00	27.68	T10
	ATOM	10644	C	LEU	A	45	-4.914	117.346	4.961	1.00	30.23	T10
	ATOM	10645	O	LEU	A	45	-5.599	118.114	4.296	1.00	26.13	T10
	ATOM	10646	N	VAL	A	46	-3.627	117.125	4.710	1.00	34.96	T10
	ATOM	10647	CA	VAL	A	46	-2.933	117.815	3.633	1.00	34.63	T10
40	ATOM	10648	CB	VAL	A	46	-1.436	117.906	3.923	1.00	40.70	T10
	ATOM	10649	CG1	VAL	A	46	-0.730	118.600	2.792	1.00	36.56	T10
	ATOM	10650	CG2	VAL	A	46	-1.211	118.656	5.203	1.00	27.41	T10
	ATOM	10651	C	VAL	A	46	-3.131	117.090	2.311	1.00	30.02	T10
	ATOM	10652	O	VAL	A	46	-2.749	115.932	2.182	1.00	28.16	T10
45	ATOM	10653	N	LYS	A	47	-3.714	117.768	1.325	1.00	26.14	T10
	ATOM	10654	CA	LYS	A	47	-3.958	117.151	0.024	1.00	28.21	T10
	ATOM	10655	CB	LYS	A	47	-5.378	117.465	-0.449	1.00	31.60	T10
	ATOM	10656	CG	LYS	A	47	-6.460	116.666	0.246	1.00	21.97	T10
	ATOM	10657	CD	LYS	A	47	-6.162	115.188	0.138	1.00	25.95	T10
50	ATOM	10658	CE	LYS	A	47	-7.344	114.331	0.542	1.00	24.99	T10
	ATOM	10659	NZ	LYS	A	47	-8.435	114.338	-0.481	1.00	31.18	T10
	ATOM	10660	C	LYS	A	47	-2.973	117.539	-1.067	1.00	29.69	T10
	ATOM	10661	O	LYS	A	47	-2.993	116.966	-2.142	1.00	29.05	T10
	ATOM	10662	N	GLU	A	48	-2.124	118.518	-0.795	1.00	28.64	T10
55	ATOM	10663	CA	GLU	A	48	-1.125	118.975	-1.761	1.00	33.18	T10
	ATOM	10664	CB	GLU	A	48	-1.598	120.214	-2.498	1.00	30.20	T10
	ATOM	10665	CG	GLU	A	48	-2.723	119.998	-3.458	1.00	34.56	T10
	ATOM	10666	CD	GLU	A	48	-3.228	121.313	-4.032	1.00	27.79	T10
	ATOM	10667	OE1	GLU	A	48	-2.383	122.190	-4.348	1.00	31.79	T10
60	ATOM	10668	OE2	GLU	A	48	-4.468	121.468	-4.171	1.00	29.56	T10
	ATOM	10669	C	GLU	A	48	0.101	119.359	-0.977	1.00	31.09	T10
	ATOM	10670	O	GLU	A	48	0.014	120.172	-0.062	1.00	30.18	T10
	ATOM	10671	N	THR	A	49	1.250	118.802	-1.324	1.00	31.40	T10
	ATOM	10672	CA	THR	A	49	2.445	119.155	-0.579	1.00	32.06	T10
65	ATOM	10673	CB	THR	A	49	3.610	118.223	-0.917	1.00	30.30	T10
	ATOM	10674	OG1	THR	A	49	4.345	118.773	-2.005	1.00	34.74	T10

	ATOM	10675	CG2	THR	A	49	3.092	116.854	-1.319	1.00	29.71	T10
	ATOM	10676	C	THR	A	49	2.825	120.605	-0.887	1.00	31.61	T10
	ATOM	10677	O	THR	A	49	2.550	121.115	-1.970	1.00	34.05	T10
	ATOM	10678	N	GLY	A	50	3.437	121.268	0.087	1.00	32.70	T10
5	ATOM	10679	CA	GLY	A	50	3.841	122.645	-0.089	1.00	34.95	T10
	ATOM	10680	C	GLY	A	50	4.161	123.278	1.252	1.00	28.08	T10
	ATOM	10681	O	GLY	A	50	4.341	122.570	2.242	1.00	33.91	T10
	ATOM	10682	N	TYR	A	51	4.240	124.608	1.281	1.00	30.63	T10
	ATOM	10683	CA	TYR	A	51	4.520	125.346	2.509	1.00	27.05	T10
10	ATOM	10684	CB	TYR	A	51	5.536	126.460	2.247	1.00	29.33	T10
	ATOM	10685	CG	TYR	A	51	6.930	125.926	2.065	1.00	37.74	T10
	ATOM	10686	CD1	TYR	A	51	7.334	125.369	0.851	1.00	31.74	T10
	ATOM	10687	CE1	TYR	A	51	8.584	124.764	0.719	1.00	33.15	T10
	ATOM	10688	CD2	TYR	A	51	7.811	125.877	3.142	1.00	33.27	T10
15	ATOM	10689	CE2	TYR	A	51	9.056	125.276	3.029	1.00	34.09	T10
	ATOM	10690	CZ	TYR	A	51	9.442	124.715	1.820	1.00	29.72	T10
	ATOM	10691	OH	TYR	A	51	10.663	124.063	1.742	1.00	29.82	T10
	ATOM	10692	C	TYR	A	51	3.236	125.933	3.084	1.00	36.67	T10
	ATOM	10693	O	TYR	A	51	2.484	126.619	2.392	1.00	30.22	T10
20	ATOM	10694	N	PHE	A	52	2.987	125.657	4.357	1.00	28.28	T10
	ATOM	10695	CA	PHE	A	52	1.785	126.144	5.003	1.00	27.68	T10
	ATOM	10696	CB	PHE	A	52	0.871	124.973	5.379	1.00	30.59	T10
	ATOM	10697	CG	PHE	A	52	0.459	124.117	4.222	1.00	35.02	T10
	ATOM	10698	CD1	PHE	A	52	1.352	123.224	3.652	1.00	33.49	T10
25	ATOM	10699	CD2	PHE	A	52	-0.833	124.190	3.711	1.00	25.06	T10
	ATOM	10700	CE1	PHE	A	52	0.966	122.416	2.590	1.00	28.41	T10
	ATOM	10701	CE2	PHE	A	52	-1.228	123.390	2.654	1.00	36.33	T10
	ATOM	10702	CZ	PHE	A	52	-0.328	122.501	2.090	1.00	31.76	T10
	ATOM	10703	C	PHE	A	52	2.047	126.956	6.264	1.00	28.96	T10
30	ATOM	10704	O	PHE	A	52	3.017	126.719	6.989	1.00	31.61	T10
	ATOM	10705	N	PHE	A	53	1.157	127.917	6.506	1.00	25.82	T10
	ATOM	10706	CA	PHE	A	53	1.194	128.757	7.694	1.00	29.87	T10
	ATOM	10707	CB	PHE	A	53	0.730	130.173	7.373	1.00	31.97	T10
	ATOM	10708	CG	PHE	A	53	0.544	131.032	8.581	1.00	30.88	T10
35	ATOM	10709	CD1	PHE	A	53	1.635	131.424	9.347	1.00	32.10	T10
	ATOM	10710	CD2	PHE	A	53	-0.724	131.438	8.969	1.00	36.25	T10
	ATOM	10711	CE1	PHE	A	53	1.458	132.209	10.485	1.00	18.11	T10
	ATOM	10712	CE2	PHE	A	53	-0.907	132.221	10.102	1.00	31.18	T10
	ATOM	10713	CZ	PHE	A	53	0.181	132.607	10.861	1.00	31.80	T10
40	ATOM	10714	C	PHE	A	53	0.166	128.083	8.575	1.00	30.70	T10
	ATOM	10715	O	PHE	A	53	-0.970	127.912	8.164	1.00	28.28	T10
	ATOM	10716	N	ILE	A	54	0.560	127.691	9.777	1.00	29.25	T10
	ATOM	10717	CA	ILE	A	54	-0.351	126.999	10.677	1.00	31.17	T10
	ATOM	10718	CB	ILE	A	54	0.166	125.579	10.955	1.00	30.74	T10
45	ATOM	10719	CG2	ILE	A	54	-0.873	124.779	11.710	1.00	32.02	T10
	ATOM	10720	CG1	ILE	A	54	0.489	124.897	9.627	1.00	29.01	T10
	ATOM	10721	CD1	ILE	A	54	1.394	123.727	9.758	1.00	32.78	T10
	ATOM	10722	C	ILE	A	54	-0.504	127.754	11.985	1.00	33.82	T10
	ATOM	10723	O	ILE	A	54	0.479	128.140	12.603	1.00	35.91	T10
50	ATOM	10724	N	TYR	A	55	-1.744	127.963	12.403	1.00	37.66	T10
	ATOM	10725	CA	TYR	A	55	-2.003	128.687	13.631	1.00	34.07	T10
	ATOM	10726	CB	TYR	A	55	-2.519	130.087	13.299	1.00	35.42	T10
	ATOM	10727	CG	TYR	A	55	-3.754	130.109	12.431	1.00	30.62	T10
	ATOM	10728	CD1	TYR	A	55	-5.022	130.223	12.989	1.00	26.79	T10
55	ATOM	10729	CE1	TYR	A	55	-6.155	130.213	12.197	1.00	27.62	T10
	ATOM	10730	CD2	TYR	A	55	-3.653	129.989	11.052	1.00	33.13	T10
	ATOM	10731	CE2	TYR	A	55	-4.779	129.977	10.247	1.00	24.00	T10
	ATOM	10732	CZ	TYR	A	55	-6.027	130.087	10.825	1.00	23.37	T10
	ATOM	10733	OH	TYR	A	55	-7.141	130.055	10.024	1.00	30.88	T10
60	ATOM	10734	C	TYR	A	55	-2.990	127.969	14.527	1.00	30.73	T10
	ATOM	10735	O	TYR	A	55	-3.577	126.972	14.139	1.00	29.58	T10
	ATOM	10736	N	GLY	A	56	-3.161	128.474	15.740	1.00	33.65	T10
	ATOM	10737	CA	GLY	A	56	-4.095	127.857	16.659	1.00	31.07	T10
	ATOM	10738	C	GLY	A	56	-4.108	128.541	18.006	1.00	29.23	T10
65	ATOM	10739	O	GLY	A	56	-3.057	128.917	18.516	1.00	27.34	T10
	ATOM	10740	N	GLN	A	57	-5.302	128.707	18.570	1.00	31.87	T10

	ATOM	10741	CA	GLN	A	57	-5.471	129.339	19.867	1.00	34.90	T10
	ATOM	10742	CB	GLN	A	57	-6.022	130.762	19.713	1.00	24.25	T10
	ATOM	10743	CG	GLN	A	57	-6.374	131.420	21.059	1.00	21.84	T10
	ATOM	10744	CD	GLN	A	57	-6.795	132.873	20.945	1.00	27.34	T10
5	ATOM	10745	OE1	GLN	A	57	-5.985	133.741	20.643	1.00	34.52	T10
	ATOM	10746	NE2	GLN	A	57	-8.069	133.141	21.197	1.00	24.38	T10
	ATOM	10747	C	GLN	A	57	-6.425	128.524	20.723	1.00	29.89	T10
	ATOM	10748	O	GLN	A	57	-7.348	127.900	20.209	1.00	23.65	T10
	ATOM	10749	N	VAL	A	58	-6.191	128.540	22.033	1.00	30.22	T10
10	ATOM	10750	CA	VAL	A	58	-7.014	127.821	23.005	1.00	26.74	T10
	ATOM	10751	CB	VAL	A	58	-6.345	126.488	23.440	1.00	36.33	T10
	ATOM	10752	CG1	VAL	A	58	-7.071	125.899	24.625	1.00	23.49	T10
	ATOM	10753	CG2	VAL	A	58	-6.348	125.503	22.294	1.00	34.21	T10
	ATOM	10754	C	VAL	A	58	-7.181	128.694	24.245	1.00	34.64	T10
15	ATOM	10755	O	VAL	A	58	-6.242	129.348	24.667	1.00	29.05	T10
	ATOM	10756	N	LEU	A	59	-8.377	128.713	24.819	1.00	29.37	T10
	ATOM	10757	CA	LEU	A	59	-8.630	129.491	26.028	1.00	32.88	T10
	ATOM	10758	CB	LEU	A	59	-10.003	130.158	25.956	1.00	35.69	T10
	ATOM	10759	CG	LEU	A	59	-10.199	131.409	26.823	1.00	34.16	T10
20	ATOM	10760	CD1	LEU	A	59	-11.677	131.670	26.988	1.00	32.02	T10
	ATOM	10761	CD2	LEU	A	59	-9.564	131.236	28.196	1.00	26.90	T10
	ATOM	10762	C	LEU	A	59	-8.590	128.535	27.234	1.00	28.01	T10
	ATOM	10763	O	LEU	A	59	-9.451	127.657	27.372	1.00	41.03	T10
	ATOM	10764	N	TYR	A	60	-7.601	128.706	28.105	1.00	32.12	T10
25	ATOM	10765	CA	TYR	A	60	-7.476	127.839	29.270	1.00	32.90	T10
	ATOM	10766	CB	TYR	A	60	-6.010	127.606	29.595	1.00	29.98	T10
	ATOM	10767	CG	TYR	A	60	-5.289	126.960	28.460	1.00	34.71	T10
	ATOM	10768	CD1	TYR	A	60	-4.417	127.687	27.660	1.00	32.35	T10
	ATOM	10769	CE1	TYR	A	60	-3.805	127.108	26.560	1.00	29.17	T10
30	ATOM	10770	CD2	TYR	A	60	-5.528	125.634	28.141	1.00	28.68	T10
	ATOM	10771	CE2	TYR	A	60	-4.930	125.044	27.056	1.00	29.60	T10
	ATOM	10772	CZ	TYR	A	60	-4.072	125.785	26.265	1.00	31.54	T10
	ATOM	10773	OH	TYR	A	60	-3.502	125.202	25.157	1.00	31.93	T10
	ATOM	10774	C	TYR	A	60	-8.177	128.365	30.509	1.00	33.37	T10
35	ATOM	10775	O	TYR	A	60	-7.939	129.494	30.949	1.00	31.22	T10
	ATOM	10776	N	THR	A	61	-9.036	127.532	31.082	1.00	29.33	T10
	ATOM	10777	CA	THR	A	61	-9.770	127.913	32.280	1.00	29.85	T10
	ATOM	10778	CB	THR	A	61	-11.284	127.905	32.031	1.00	36.97	T10
	ATOM	10779	OG1	THR	A	61	-11.682	126.631	31.512	1.00	28.42	T10
40	ATOM	10780	CG2	THR	A	61	-11.650	128.989	31.034	1.00	28.87	T10
	ATOM	10781	C	THR	A	61	-9.424	126.927	33.376	1.00	34.64	T10
	ATOM	10782	O	THR	A	61	-10.100	126.833	34.393	1.00	31.02	T10
	ATOM	10783	N	ASP	A	62	-8.346	126.190	33.145	1.00	34.74	T10
	ATOM	10784	CA	ASP	A	62	-7.843	125.208	34.093	1.00	29.77	T10
45	ATOM	10785	CB	ASP	A	62	-7.075	124.134	33.316	1.00	33.01	T10
	ATOM	10786	CG	ASP	A	62	-6.786	122.895	34.145	1.00	29.58	T10
	ATOM	10787	OD1	ASP	A	62	-7.074	121.774	33.641	1.00	35.11	T10
	ATOM	10788	OD2	ASP	A	62	-6.264	123.045	35.282	1.00	29.23	T10
	ATOM	10789	C	ASP	A	62	-6.914	125.953	35.061	1.00	33.31	T10
50	ATOM	10790	O	ASP	A	62	-6.268	126.926	34.684	1.00	37.22	T10
	ATOM	10791	N	LYS	A	63	-6.837	125.516	36.309	1.00	30.78	T10
	ATOM	10792	CA	LYS	A	63	-5.970	126.214	37.254	1.00	34.16	T10
	ATOM	10793	CB	LYS	A	63	-6.743	126.549	38.528	1.00	37.32	T10
	ATOM	10794	CG	LYS	A	63	-7.294	125.329	39.245	1.00	27.38	T10
55	ATOM	10795	CD	LYS	A	63	-8.018	125.730	40.537	1.00	24.12	T10
	ATOM	10796	CE	LYS	A	63	-9.223	126.665	40.255	1.00	25.40	T10
	ATOM	10797	NZ	LYS	A	63	-9.954	127.108	41.505	1.00	31.39	T10
	ATOM	10798	C	LYS	A	63	-4.697	125.458	37.629	1.00	30.81	T10
	ATOM	10799	O	LYS	A	63	-4.093	125.738	38.673	1.00	36.10	T10
60	ATOM	10800	N	THR	A	64	-4.263	124.529	36.783	1.00	31.09	T10
	ATOM	10801	CA	THR	A	64	-3.071	123.768	37.125	1.00	30.94	T10
	ATOM	10802	CB	THR	A	64	-3.123	122.324	36.531	1.00	28.86	T10
	ATOM	10803	OG1	THR	A	64	-3.412	122.371	35.125	1.00	24.92	T10
	ATOM	10804	CG2	THR	A	64	-4.195	121.499	37.257	1.00	29.00	T10
65	ATOM	10805	C	THR	A	64	-1.729	124.416	36.790	1.00	30.74	T10
	ATOM	10806	O	THR	A	64	-0.940	123.866	36.028	1.00	32.41	T10

	ATOM	10807	N	TYR	A	65	-1.483	125.585	37.374	1.00	33.59	T10
	ATOM	10808	CA	TYR	A	65	-0.224	126.323	37.213	1.00	31.09	T10
	ATOM	10809	CB	TYR	A	65	0.839	125.713	38.147	1.00	25.53	T10
	ATOM	10810	CG	TYR	A	65	1.867	124.837	37.461	1.00	29.11	T10
5	ATOM	10811	CD1	TYR	A	65	3.070	125.373	36.978	1.00	26.16	T10
	ATOM	10812	CE1	TYR	A	65	4.017	124.565	36.324	1.00	29.15	T10
	ATOM	10813	CD2	TYR	A	65	1.630	123.477	37.276	1.00	29.46	T10
	ATOM	10814	CE2	TYR	A	65	2.563	122.657	36.624	1.00	32.62	T10
	ATOM	10815	CZ	TYR	A	65	3.753	123.202	36.150	1.00	28.14	T10
10	ATOM	10816	OH	TYR	A	65	4.665	122.374	35.517	1.00	31.07	T10
	ATOM	10817	C	TYR	A	65	0.391	126.510	35.809	1.00	28.10	T10
	ATOM	10818	O	TYR	A	65	1.167	127.449	35.590	1.00	27.93	T10
	ATOM	10819	N	ALA	A	66	0.076	125.635	34.864	1.00	35.27	T10
	ATOM	10820	CA	ALA	A	66	0.642	125.773	33.533	1.00	30.00	T10
15	ATOM	10821	CB	ALA	A	66	2.092	125.335	33.549	1.00	26.30	T10
	ATOM	10822	C	ALA	A	66	-0.136	124.966	32.508	1.00	32.86	T10
	ATOM	10823	O	ALA	A	66	-0.222	123.742	32.596	1.00	27.73	T10
	ATOM	10824	N	MET	A	67	-0.710	125.657	31.532	1.00	27.78	T10
	ATOM	10825	CA	MET	A	67	-1.464	124.993	30.486	1.00	32.61	T10
20	ATOM	10826	CB	MET	A	67	-2.920	125.454	30.508	1.00	27.81	T10
	ATOM	10827	CG	MET	A	67	-3.699	124.963	31.717	1.00	26.21	T10
	ATOM	10828	SD	MET	A	67	-3.733	123.170	31.799	1.00	36.96	T10
	ATOM	10829	CE	MET	A	67	-5.037	122.777	30.593	1.00	28.32	T10
	ATOM	10830	C	MET	A	67	-0.833	125.324	29.148	1.00	32.15	T10
25	ATOM	10831	O	MET	A	67	0.021	126.209	29.064	1.00	28.69	T10
	ATOM	10832	N	GLY	A	68	-1.246	124.611	28.105	1.00	23.29	T10
	ATOM	10833	CA	GLY	A	68	-0.702	124.860	26.779	1.00	30.60	T10
	ATOM	10834	C	GLY	A	68	-1.099	123.789	25.782	1.00	31.94	T10
	ATOM	10835	O	GLY	A	68	-1.658	122.758	26.165	1.00	33.15	T10
30	ATOM	10836	N	HIS	A	69	-0.833	124.032	24.502	1.00	32.15	T10
	ATOM	10837	CA	HIS	A	69	-1.165	123.058	23.472	1.00	27.93	T10
	ATOM	10838	CB	HIS	A	69	-2.465	123.436	22.739	1.00	25.35	T10
	ATOM	10839	CG	HIS	A	69	-2.480	124.821	22.166	1.00	33.24	T10
	ATOM	10840	CD2	HIS	A	69	-2.354	125.259	20.891	1.00	24.95	T10
35	ATOM	10841	ND1	HIS	A	69	-2.683	125.948	22.935	1.00	30.04	T10
	ATOM	10842	CE1	HIS	A	69	-2.686	127.016	22.161	1.00	30.03	T10
	ATOM	10843	NE2	HIS	A	69	-2.488	126.626	20.914	1.00	36.40	T10
	ATOM	10844	C	HIS	A	69	-0.043	122.860	22.468	1.00	33.22	T10
	ATOM	10845	O	HIS	A	69	0.921	123.629	22.430	1.00	24.17	T10
40	ATOM	10846	N	LEU	A	70	-0.177	121.814	21.661	1.00	30.17	T10
	ATOM	10847	CA	LEU	A	70	0.817	121.483	20.651	1.00	36.30	T10
	ATOM	10848	CB	LEU	A	70	1.434	120.126	20.953	1.00	24.98	T10
	ATOM	10849	CG	LEU	A	70	1.743	119.778	22.397	1.00	34.87	T10
	ATOM	10850	CD1	LEU	A	70	2.142	118.319	22.461	1.00	22.02	T10
45	ATOM	10851	CD2	LEU	A	70	2.844	120.675	22.924	1.00	25.51	T10
	ATOM	10852	C	LEU	A	70	0.176	121.384	19.279	1.00	31.60	T10
	ATOM	10853	O	LEU	A	70	-0.962	120.941	19.151	1.00	35.46	T10
	ATOM	10854	N	ILE	A	71	0.905	121.798	18.256	1.00	38.59	T10
	ATOM	10855	CA	ILE	A	71	0.412	121.674	16.900	1.00	35.27	T10
50	ATOM	10856	CB	ILE	A	71	0.553	122.987	16.116	1.00	38.38	T10
	ATOM	10857	CG2	ILE	A	71	0.322	122.744	14.640	1.00	32.15	T10
	ATOM	10858	CG1	ILE	A	71	-0.462	124.002	16.639	1.00	25.97	T10
	ATOM	10859	CD1	ILE	A	71	-0.393	125.337	15.944	1.00	31.08	T10
	ATOM	10860	C	ILE	A	71	1.314	120.600	16.317	1.00	28.46	T10
55	ATOM	10861	O	ILE	A	71	2.493	120.828	16.089	1.00	31.14	T10
	ATOM	10862	N	GLN	A	72	0.765	119.419	16.078	1.00	32.00	T10
	ATOM	10863	CA	GLN	A	72	1.578	118.322	15.576	1.00	30.22	T10
	ATOM	10864	CB	GLN	A	72	1.407	117.122	16.494	1.00	27.53	T10
	ATOM	10865	CG	GLN	A	72	1.493	117.509	17.947	1.00	31.87	T10
60	ATOM	10866	CD	GLN	A	72	1.527	116.314	18.853	1.00	23.52	T10
	ATOM	10867	OE1	GLN	A	72	0.645	115.450	18.804	1.00	38.01	T10
	ATOM	10868	NE2	GLN	A	72	2.551	116.252	19.697	1.00	30.28	T10
	ATOM	10869	C	GLN	A	72	1.344	117.894	14.148	1.00	24.23	T10
	ATOM	10870	O	GLN	A	72	0.281	118.116	13.575	1.00	32.68	T10
65	ATOM	10871	N	ARG	A	73	2.359	117.247	13.596	1.00	31.55	T10
	ATOM	10872	CA	ARG	A	73	2.332	116.760	12.226	1.00	28.05	T10

	ATOM	10873	CB	ARG	A	73	3.412	117.482	11.424	1.00	24.47	T10
	ATOM	10874	CG	ARG	A	73	3.491	117.042	9.997	1.00	32.51	T10
	ATOM	10875	CD	ARG	A	73	4.859	117.303	9.433	1.00	24.02	T10
	ATOM	10876	NE	ARG	A	73	4.958	116.858	8.047	1.00	31.03	T10
5	ATOM	10877	CZ	ARG	A	73	6.103	116.703	7.398	1.00	30.31	T10
	ATOM	10878	NH1	ARG	A	73	7.253	116.951	8.005	1.00	36.23	T10
	ATOM	10879	NH2	ARG	A	73	6.093	116.307	6.141	1.00	36.36	T10
	ATOM	10880	C	ARG	A	73	2.577	115.245	12.163	1.00	32.43	T10
	ATOM	10881	O	ARG	A	73	3.498	114.734	12.810	1.00	28.36	T10
10	ATOM	10882	N	LYS	A	74	1.751	114.536	11.395	1.00	29.13	T10
	ATOM	10883	CA	LYS	A	74	1.906	113.094	11.234	1.00	31.64	T10
	ATOM	10884	CB	LYS	A	74	0.571	112.366	11.385	1.00	31.72	T10
	ATOM	10885	CG	LYS	A	74	0.000	112.381	12.799	1.00	23.84	T10
	ATOM	10886	CD	LYS	A	74	-1.390	111.712	12.878	1.00	28.22	T10
15	ATOM	10887	CE	LYS	A	74	-1.967	111.791	14.309	1.00	30.43	T10
	ATOM	10888	NZ	LYS	A	74	-3.305	111.122	14.460	1.00	29.86	T10
	ATOM	10889	C	LYS	A	74	2.442	112.828	9.848	1.00	33.44	T10
	ATOM	10890	O	LYS	A	74	1.685	112.855	8.878	1.00	24.55	T10
	ATOM	10891	N	LYS	A	75	3.743	112.570	9.756	1.00	24.58	T10
20	ATOM	10892	CA	LYS	A	75	4.396	112.302	8.474	1.00	33.67	T10
	ATOM	10893	CB	LYS	A	75	5.894	112.062	8.679	1.00	31.30	T10
	ATOM	10894	CG	LYS	A	75	6.670	113.204	9.290	1.00	37.43	T10
	ATOM	10895	CD	LYS	A	75	8.137	112.822	9.409	1.00	34.87	T10
	ATOM	10896	CE	LYS	A	75	8.947	113.957	10.035	1.00	26.56	T10
25	ATOM	10897	NZ	LYS	A	75	10.378	113.609	10.293	1.00	38.00	T10
	ATOM	10898	C	LYS	A	75	3.809	111.078	7.781	1.00	29.72	T10
	ATOM	10899	O	LYS	A	75	3.595	110.055	8.416	1.00	34.31	T10
	ATOM	10900	N	VAL	A	76	3.562	111.184	6.480	1.00	33.33	T10
	ATOM	10901	CA	VAL	A	76	3.025	110.059	5.715	1.00	29.12	T10
30	ATOM	10902	CB	VAL	A	76	2.495	110.463	4.335	1.00	28.91	T10
	ATOM	10903	CG1	VAL	A	76	1.617	109.379	3.797	1.00	34.90	T10
	ATOM	10904	CG2	VAL	A	76	1.775	111.756	4.410	1.00	26.56	T10
	ATOM	10905	C	VAL	A	76	4.188	109.134	5.430	1.00	28.43	T10
	ATOM	10906	O	VAL	A	76	4.054	107.910	5.481	1.00	29.35	T10
35	ATOM	10907	N	HIS	A	77	5.323	109.744	5.099	1.00	28.89	T10
	ATOM	10908	CA	HIS	A	77	6.527	109.009	4.789	1.00	31.13	T10
	ATOM	10909	CB	HIS	A	77	7.156	109.573	3.525	1.00	26.54	T10
	ATOM	10910	CG	HIS	A	77	6.250	109.527	2.335	1.00	33.89	T10
	ATOM	10911	CD2	HIS	A	77	6.276	110.214	1.168	1.00	24.12	T10
40	ATOM	10912	ND1	HIS	A	77	5.202	108.634	2.233	1.00	25.32	T10
	ATOM	10913	CE1	HIS	A	77	4.626	108.770	1.053	1.00	29.31	T10
	ATOM	10914	NE2	HIS	A	77	5.259	109.721	0.387	1.00	28.38	T10
	ATOM	10915	C	HIS	A	77	7.479	109.111	5.963	1.00	30.01	T10
	ATOM	10916	O	HIS	A	77	7.517	110.138	6.638	1.00	30.01	T10
45	ATOM	10917	N	VAL	A	78	8.259	108.058	6.201	1.00	35.04	T10
	ATOM	10918	CA	VAL	A	78	9.154	108.076	7.336	1.00	27.97	T10
	ATOM	10919	CB	VAL	A	78	8.625	107.113	8.412	1.00	34.94	T10
	ATOM	10920	CG1	VAL	A	78	9.445	107.227	9.684	1.00	31.06	T10
	ATOM	10921	CG2	VAL	A	78	7.183	107.468	8.732	1.00	19.31	T10
50	ATOM	10922	C	VAL	A	78	10.666	107.883	7.140	1.00	26.18	T10
	ATOM	10923	O	VAL	A	78	11.427	108.820	7.386	1.00	35.41	T10
	ATOM	10924	N	PHE	A	79	11.134	106.718	6.717	1.00	34.57	T10
	ATOM	10925	CA	PHE	A	79	12.596	106.522	6.563	1.00	34.58	T10
	ATOM	10926	CB	PHE	A	79	13.279	107.717	5.880	1.00	29.60	T10
55	ATOM	10927	CG	PHE	A	79	12.656	108.127	4.587	1.00	29.05	T10
	ATOM	10928	CD1	PHE	A	79	11.812	109.227	4.532	1.00	29.62	T10
	ATOM	10929	CD2	PHE	A	79	12.912	107.418	3.421	1.00	32.58	T10
	ATOM	10930	CE1	PHE	A	79	11.230	109.620	3.338	1.00	34.62	T10
	ATOM	10931	CE2	PHE	A	79	12.336	107.799	2.223	1.00	31.02	T10
60	ATOM	10932	CZ	PHE	A	79	11.492	108.905	2.181	1.00	25.21	T10
	ATOM	10933	C	PHE	A	79	13.369	106.286	7.876	1.00	28.95	T10
	ATOM	10934	O	PHE	A	79	13.347	107.112	8.796	1.00	29.66	T10
	ATOM	10935	N	GLY	A	80	14.072	105.162	7.934	1.00	31.95	T10
	ATOM	10936	CA	GLY	A	80	14.872	104.833	9.100	1.00	28.66	T10
65	ATOM	10937	C	GLY	A	80	14.229	105.006	10.459	1.00	28.15	T10
	ATOM	10938	O	GLY	A	80	13.119	104.534	10.704	1.00	29.04	T10

	ATOM	10939	N	ASP	A	81	14.940	105.692	11.349	1.00	29.85	T10
	ATOM	10940	CA	ASP	A	81	14.474	105.910	12.713	1.00	35.77	T10
	ATOM	10941	CB	ASP	A	81	15.634	105.722	13.700	1.00	26.25	T10
	ATOM	10942	CG	ASP	A	81	16.724	106.773	13.541	1.00	28.48	T10
5	ATOM	10943	OD1	ASP	A	81	16.767	107.449	12.493	1.00	29.59	T10
	ATOM	10944	OD2	ASP	A	81	17.554	106.920	14.462	1.00	31.78	T10
	ATOM	10945	C	ASP	A	81	13.813	107.259	12.944	1.00	26.70	T10
	ATOM	10946	O	ASP	A	81	13.858	107.805	14.046	1.00	35.79	T10
	ATOM	10947	N	GLU	A	82	13.199	107.808	11.906	1.00	27.73	T10
10	ATOM	10948	CA	GLU	A	82	12.514	109.083	12.061	1.00	30.63	T10
	ATOM	10949	CB	GLU	A	82	12.022	109.619	10.723	1.00	41.11	T10
	ATOM	10950	CG	GLU	A	82	13.016	110.292	9.848	1.00	26.35	T10
	ATOM	10951	CD	GLU	A	82	12.334	111.305	8.948	1.00	34.32	T10
	ATOM	10952	OE1	GLU	A	82	11.260	110.989	8.394	1.00	40.02	T10
15	ATOM	10953	OE2	GLU	A	82	12.862	112.427	8.791	1.00	23.62	T10
	ATOM	10954	C	GLU	A	82	11.279	108.823	12.898	1.00	28.44	T10
	ATOM	10955	O	GLU	A	82	10.739	107.721	12.859	1.00	25.30	T10
	ATOM	10956	N	LEU	A	83	10.826	109.819	13.651	1.00	27.80	T10
	ATOM	10957	CA	LEU	A	83	9.591	109.663	14.415	1.00	30.64	T10
20	ATOM	10958	CB	LEU	A	83	9.601	110.485	15.700	1.00	32.70	T10
	ATOM	10959	CG	LEU	A	83	10.557	110.020	16.792	1.00	30.47	T10
	ATOM	10960	CD1	LEU	A	83	12.001	110.157	16.314	1.00	38.40	T10
	ATOM	10961	CD2	LEU	A	83	10.340	110.852	18.028	1.00	29.28	T10
	ATOM	10962	C	LEU	A	83	8.575	110.236	13.460	1.00	29.19	T10
25	ATOM	10963	O	LEU	A	83	8.785	111.328	12.927	1.00	28.60	T10
	ATOM	10964	N	SER	A	84	7.492	109.507	13.219	1.00	32.23	T10
	ATOM	10965	CA	SER	A	84	6.479	109.977	12.283	1.00	32.83	T10
	ATOM	10966	CB	SER	A	84	5.646	108.802	11.763	1.00	33.75	T10
	ATOM	10967	OG	SER	A	84	5.223	107.977	12.826	1.00	30.69	T10
30	ATOM	10968	C	SER	A	84	5.570	111.053	12.855	1.00	31.14	T10
	ATOM	10969	O	SER	A	84	4.726	111.595	12.149	1.00	21.10	T10
	ATOM	10970	N	LEU	A	85	5.751	111.372	14.130	1.00	28.85	T10
	ATOM	10971	CA	LEU	A	85	4.946	112.401	14.773	1.00	32.29	T10
	ATOM	10972	CB	LEU	A	85	4.207	111.807	15.974	1.00	34.59	T10
35	ATOM	10973	CG	LEU	A	85	3.016	112.548	16.594	1.00	27.50	T10
	ATOM	10974	CD1	LEU	A	85	3.482	113.786	17.336	1.00	28.74	T10
	ATOM	10975	CD2	LEU	A	85	2.019	112.904	15.506	1.00	28.12	T10
	ATOM	10976	C	LEU	A	85	5.871	113.524	15.222	1.00	29.07	T10
	ATOM	10977	O	LEU	A	85	6.629	113.371	16.180	1.00	32.75	T10
40	ATOM	10978	N	VAL	A	86	5.821	114.649	14.516	1.00	29.40	T10
	ATOM	10979	CA	VAL	A	86	6.657	115.797	14.857	1.00	29.35	T10
	ATOM	10980	CB	VAL	A	86	7.362	116.385	13.632	1.00	28.22	T10
	ATOM	10981	CG1	VAL	A	86	8.126	117.633	14.033	1.00	26.67	T10
	ATOM	10982	CG2	VAL	A	86	8.303	115.356	13.026	1.00	32.04	T10
45	ATOM	10983	C	VAL	A	86	5.773	116.877	15.415	1.00	28.81	T10
	ATOM	10984	O	VAL	A	86	4.682	117.105	14.904	1.00	23.31	T10
	ATOM	10985	N	THR	A	87	6.232	117.541	16.465	1.00	30.37	T10
	ATOM	10986	CA	THR	A	87	5.440	118.609	17.034	1.00	27.63	T10
	ATOM	10987	CB	THR	A	87	5.275	118.454	18.572	1.00	32.63	T10
50	ATOM	10988	OG1	THR	A	87	5.872	119.570	19.239	1.00	31.97	T10
	ATOM	10989	CG2	THR	A	87	5.907	117.152	19.052	1.00	23.99	T10
	ATOM	10990	C	THR	A	87	6.093	119.932	16.683	1.00	28.61	T10
	ATOM	10991	O	THR	A	87	7.229	120.197	17.039	1.00	31.61	T10
	ATOM	10992	N	LEU	A	88	5.361	120.744	15.932	1.00	25.47	T10
55	ATOM	10993	CA	LEU	A	88	5.807	122.064	15.502	1.00	23.75	T10
	ATOM	10994	CB	LEU	A	88	5.239	122.368	14.115	1.00	33.24	T10
	ATOM	10995	CG	LEU	A	88	5.436	121.487	12.878	1.00	25.91	T10
	ATOM	10996	CD1	LEU	A	88	5.607	120.058	13.241	1.00	25.26	T10
	ATOM	10997	CD2	LEU	A	88	4.235	121.637	11.988	1.00	23.81	T10
60	ATOM	10998	C	LEU	A	88	5.219	123.087	16.476	1.00	30.25	T10
	ATOM	10999	O	LEU	A	88	4.112	122.911	16.995	1.00	36.11	T10
	ATOM	11000	N	PHE	A	89	5.942	124.151	16.766	1.00	33.14	T10
	ATOM	11001	CA	PHE	A	89	5.352	125.192	17.620	1.00	31.69	T10
	ATOM	11002	CB	PHE	A	89	4.076	125.708	16.939	1.00	28.97	T10
65	ATOM	11003	CG	PHE	A	89	4.244	125.896	15.459	1.00	24.24	T10
	ATOM	11004	CD1	PHE	A	89	3.201	125.641	14.584	1.00	20.28	T10

	ATOM	11005	CD2	PHE	A	89	5.502	126.233	14.927	1.00	30.27	T10
	ATOM	11006	CE1	PHE	A	89	3.412	125.706	13.190	1.00	29.50	T10
	ATOM	11007	CE2	PHE	A	89	5.718	126.300	13.544	1.00	35.65	T10
	ATOM	11008	CZ	PHE	A	89	4.675	126.033	12.676	1.00	36.32	T10
5	ATOM	11009	C	PHE	A	89	5.075	124.895	19.095	1.00	32.35	T10
	ATOM	11010	O	PHE	A	89	6.014	124.871	19.895	1.00	32.56	T10
	ATOM	11011	N	ARG	A	90	3.814	124.704	19.485	1.00	32.97	T10
	ATOM	11012	CA	ARG	A	90	3.532	124.469	20.913	1.00	27.29	T10
	ATOM	11013	CB	ARG	A	90	4.385	123.305	21.427	1.00	26.81	T10
10	ATOM	11014	CG	ARG	A	90	4.708	123.383	22.914	1.00	27.90	T10
	ATOM	11015	CD	ARG	A	90	5.853	122.464	23.272	1.00	26.61	T10
	ATOM	11016	NE	ARG	A	90	6.285	122.699	24.642	1.00	37.90	T10
	ATOM	11017	CZ	ARG	A	90	7.550	122.644	25.043	1.00	26.86	T10
	ATOM	11018	NH1	ARG	A	90	8.519	122.355	24.178	1.00	33.74	T10
15	ATOM	11019	NH2	ARG	A	90	7.851	122.909	26.307	1.00	31.87	T10
	ATOM	11020	C	ARG	A	90	3.781	125.696	21.838	1.00	31.39	T10
	ATOM	11021	O	ARG	A	90	4.862	126.287	21.830	1.00	26.97	T10
	ATOM	11022	N	CYS	A	91	2.791	126.047	22.656	1.00	32.55	T10
	ATOM	11023	CA	CYS	A	91	2.916	127.180	23.573	1.00	34.14	T10
20	ATOM	11024	CB	CYS	A	91	2.049	128.334	23.090	1.00	31.09	T10
	ATOM	11025	SG	CYS	A	91	0.338	127.865	22.846	1.00	32.74	T10
	ATOM	11026	C	CYS	A	91	2.530	126.810	25.009	1.00	22.01	T10
	ATOM	11027	O	CYS	A	91	1.941	125.757	25.246	1.00	32.91	T10
	ATOM	11028	N	ILE	A	92	2.859	127.681	25.962	1.00	31.59	T10
25	ATOM	11029	CA	ILE	A	92	2.574	127.437	27.378	1.00	33.20	T10
	ATOM	11030	CB	ILE	A	92	3.838	126.983	28.119	1.00	32.04	T10
	ATOM	11031	CG2	ILE	A	92	3.514	126.673	29.565	1.00	32.65	T10
	ATOM	11032	CG1	ILE	A	92	4.419	125.738	27.444	1.00	27.80	T10
	ATOM	11033	CD1	ILE	A	92	5.840	125.389	27.914	1.00	31.63	T10
30	ATOM	11034	C	ILE	A	92	2.101	128.715	28.052	1.00	34.65	T10
	ATOM	11035	O	ILE	A	92	2.414	129.800	27.600	1.00	21.69	T10
	ATOM	11036	N	GLN	A	93	1.345	128.585	29.135	1.00	28.69	T10
	ATOM	11037	CA	GLN	A	93	0.863	129.748	29.874	1.00	29.56	T10
	ATOM	11038	CB	GLN	A	93	-0.445	130.260	29.274	1.00	30.99	T10
35	ATOM	11039	CG	GLN	A	93	-0.273	131.247	28.122	1.00	31.44	T10
	ATOM	11040	CD	GLN	A	93	-0.380	132.709	28.570	1.00	23.30	T10
	ATOM	11041	OE1	GLN	A	93	0.540	133.267	29.186	1.00	26.06	T10
	ATOM	11042	NE2	GLN	A	93	-1.516	133.331	28.270	1.00	34.51	T10
	ATOM	11043	C	GLN	A	93	0.649	129.411	31.344	1.00	27.05	T10
40	ATOM	11044	O	GLN	A	93	-0.124	128.500	31.679	1.00	31.53	T10
	ATOM	11045	N	ASN	A	94	1.351	130.124	32.224	1.00	31.19	T10
	ATOM	11046	CA	ASN	A	94	1.189	129.891	33.649	1.00	33.32	T10
	ATOM	11047	CB	ASN	A	94	2.126	130.793	34.451	1.00	38.83	T10
	ATOM	11048	CG	ASN	A	94	3.536	130.248	34.520	1.00	25.86	T10
45	ATOM	11049	OD1	ASN	A	94	3.747	129.101	34.924	1.00	38.27	T10
	ATOM	11050	ND2	ASN	A	94	4.513	131.063	34.138	1.00	38.84	T10
	ATOM	11051	C	ASN	A	94	-0.265	130.225	33.968	1.00	31.12	T10
	ATOM	11052	O	ASN	A	94	-0.854	131.122	33.357	1.00	30.28	T10
	ATOM	11053	N	MET	A	95	-0.854	129.491	34.904	1.00	27.33	T10
50	ATOM	11054	CA	MET	A	95	-2.239	129.729	35.292	1.00	33.67	T10
	ATOM	11055	CB	MET	A	95	-3.062	128.441	35.164	1.00	31.57	T10
	ATOM	11056	CG	MET	A	95	-3.123	127.874	33.771	1.00	37.42	T10
	ATOM	11057	SD	MET	A	95	-3.663	129.101	32.546	1.00	27.53	T10
	ATOM	11058	CE	MET	A	95	-5.426	129.224	32.871	1.00	37.43	T10
55	ATOM	11059	C	MET	A	95	-2.328	130.228	36.734	1.00	36.81	T10
	ATOM	11060	O	MET	A	95	-1.476	129.905	37.571	1.00	31.12	T10
	ATOM	11061	N	PRO	A	96	-3.364	131.031	37.039	1.00	27.07	T10
	ATOM	11062	CD	PRO	A	96	-4.363	131.582	36.112	1.00	27.75	T10
	ATOM	11063	CA	PRO	A	96	-3.576	131.570	38.382	1.00	33.41	T10
60	ATOM	11064	CB	PRO	A	96	-4.602	132.687	38.172	1.00	32.03	T10
	ATOM	11065	CG	PRO	A	96	-4.563	132.956	36.702	1.00	23.59	T10
	ATOM	11066	C	PRO	A	96	-4.184	130.429	39.177	1.00	29.55	T10
	ATOM	11067	O	PRO	A	96	-4.240	129.285	38.708	1.00	31.40	T10
	ATOM	11068	N	GLU	A	97	-4.674	130.740	40.365	1.00	30.37	T10
65	ATOM	11069	CA	GLU	A	97	-5.267	129.709	41.190	1.00	34.04	T10
	ATOM	11070	CB	GLU	A	97	-4.572	129.679	42.542	1.00	31.10	T10

	ATOM	11071	CG	GLU	A	97	-4.849	128.420	43.310	1.00	31.32	T10
	ATOM	11072	CD	GLU	A	97	-3.578	127.876	43.938	1.00	30.60	T10
	ATOM	11073	OE1	GLU	A	97	-2.965	128.609	44.759	1.00	30.28	T10
	ATOM	11074	OE2	GLU	A	97	-3.184	126.725	43.604	1.00	31.29	T10
5	ATOM	11075	C	GLU	A	97	-6.747	129.981	41.379	1.00	36.48	T10
	ATOM	11076	O	GLU	A	97	-7.535	129.067	41.656	1.00	26.93	T10
	ATOM	11077	N	THR	A	98	-7.118	131.244	41.200	1.00	28.02	T10
	ATOM	11078	CA	THR	A	98	-8.494	131.653	41.395	1.00	28.98	T10
	ATOM	11079	CB	THR	A	98	-8.548	133.045	42.015	1.00	35.16	T10
10	ATOM	11080	OG1	THR	A	98	-7.970	133.985	41.098	1.00	36.08	T10
	ATOM	11081	CG2	THR	A	98	-7.771	133.066	43.320	1.00	31.84	T10
	ATOM	11082	C	THR	A	98	-9.375	131.656	40.144	1.00	36.10	T10
	ATOM	11083	O	THR	A	98	-10.257	130.799	40.004	1.00	29.71	T10
	ATOM	11084	N	LEU	A	99	-9.155	132.598	39.232	1.00	32.49	T10
15	ATOM	11085	CA	LEU	A	99	-10.022	132.654	38.066	1.00	37.41	T10
	ATOM	11086	CB	LEU	A	99	-10.767	133.988	38.056	1.00	21.60	T10
	ATOM	11087	CG	LEU	A	99	-11.675	134.159	39.284	1.00	28.07	T10
	ATOM	11088	CD1	LEU	A	99	-12.278	135.561	39.335	1.00	30.76	T10
	ATOM	11089	CD2	LEU	A	99	-12.774	133.101	39.226	1.00	35.63	T10
20	ATOM	11090	C	LEU	A	99	-9.329	132.422	36.738	1.00	31.61	T10
	ATOM	11091	O	LEU	A	99	-9.172	133.348	35.931	1.00	34.16	T10
	ATOM	11092	N	PRO	A	100	-8.925	131.165	36.482	1.00	30.46	T10
	ATOM	11093	CD	PRO	A	100	-9.178	129.984	37.328	1.00	25.74	T10
	ATOM	11094	CA	PRO	A	100	-8.240	130.766	35.253	1.00	35.39	T10
25	ATOM	11095	CB	PRO	A	100	-8.321	129.243	35.299	1.00	33.93	T10
	ATOM	11096	CG	PRO	A	100	-8.209	128.966	36.754	1.00	29.06	T10
	ATOM	11097	C	PRO	A	100	-8.879	131.343	33.996	1.00	27.02	T10
	ATOM	11098	O	PRO	A	100	-10.059	131.128	33.721	1.00	34.08	T10
	ATOM	11099	N	ASN	A	101	-8.079	132.073	33.237	1.00	35.05	T10
30	ATOM	11100	CA	ASN	A	101	-8.537	132.686	32.005	1.00	34.38	T10
	ATOM	11101	CB	ASN	A	101	-9.384	133.906	32.306	1.00	29.03	T10
	ATOM	11102	CG	ASN	A	101	-10.815	133.563	32.503	1.00	29.02	T10
	ATOM	11103	OD1	ASN	A	101	-11.500	133.167	31.558	1.00	28.16	T10
	ATOM	11104	ND2	ASN	A	101	-11.291	133.696	33.739	1.00	24.43	T10
35	ATOM	11105	C	ASN	A	101	-7.352	133.135	31.194	1.00	32.14	T10
	ATOM	11106	O	ASN	A	101	-6.988	134.315	31.245	1.00	31.92	T10
	ATOM	11107	N	ASN	A	102	-6.741	132.231	30.435	1.00	35.82	T10
	ATOM	11108	CA	ASN	A	102	-5.598	132.687	29.680	1.00	25.31	T10
	ATOM	11109	CB	ASN	A	102	-4.322	132.048	30.223	1.00	29.71	T10
40	ATOM	11110	CG	ASN	A	102	-3.776	132.817	31.416	1.00	29.34	T10
	ATOM	11111	OD1	ASN	A	102	-3.709	134.057	31.388	1.00	24.16	T10
	ATOM	11112	ND2	ASN	A	102	-3.393	132.099	32.469	1.00	31.55	T10
	ATOM	11113	C	ASN	A	102	-5.594	132.710	28.159	1.00	27.23	T10
	ATOM	11114	O	ASN	A	102	-5.206	133.735	27.588	1.00	28.40	T10
45	ATOM	11115	N	SER	A	103	-6.007	131.659	27.466	1.00	29.53	T10
	ATOM	11116	CA	SER	A	103	-5.968	131.785	26.009	1.00	32.96	T10
	ATOM	11117	CB	SER	A	103	-6.891	132.933	25.551	1.00	27.44	T10
	ATOM	11118	OG	SER	A	103	-6.248	133.789	24.617	1.00	34.15	T10
	ATOM	11119	C	SER	A	103	-4.530	132.044	25.507	1.00	29.68	T10
50	ATOM	11120	O	SER	A	103	-3.819	132.934	25.976	1.00	26.67	T10
	ATOM	11121	N	CYS	A	104	-4.109	131.263	24.528	1.00	29.66	T10
	ATOM	11122	CA	CYS	A	104	-2.773	131.412	24.010	1.00	30.13	T10
	ATOM	11123	CB	CYS	A	104	-1.852	130.479	24.775	1.00	34.16	T10
	ATOM	11124	SG	CYS	A	104	-0.122	130.756	24.491	1.00	31.55	T10
55	ATOM	11125	C	CYS	A	104	-2.754	131.086	22.525	1.00	27.91	T10
	ATOM	11126	O	CYS	A	104	-3.217	130.028	22.104	1.00	30.72	T10
	ATOM	11127	N	TYR	A	105	-2.226	132.014	21.736	1.00	31.75	T10
	ATOM	11128	CA	TYR	A	105	-2.133	131.856	20.295	1.00	28.50	T10
	ATOM	11129	CB	TYR	A	105	-2.608	133.132	19.606	1.00	24.19	T10
60	ATOM	11130	CG	TYR	A	105	-2.459	133.133	18.103	1.00	31.11	T10
	ATOM	11131	CD1	TYR	A	105	-3.509	132.754	17.282	1.00	28.92	T10
	ATOM	11132	CE1	TYR	A	105	-3.384	132.769	15.895	1.00	37.02	T10
	ATOM	11133	CD2	TYR	A	105	-1.266	133.527	17.499	1.00	30.01	T10
	ATOM	11134	CE2	TYR	A	105	-1.130	133.542	16.112	1.00	36.67	T10
65	ATOM	11135	CZ	TYR	A	105	-2.195	133.163	15.320	1.00	25.02	T10
	ATOM	11136	OH	TYR	A	105	-2.073	133.196	13.953	1.00	32.32	T10

	ATOM	11137	C	TYR	A	105	-0.683	131.586	19.910	1.00	33.12	T10
	ATOM	11138	O	TYR	A	105	0.244	132.084	20.548	1.00	29.86	T10
	ATOM	11139	N	SER	A	106	-0.485	130.793	18.866	1.00	24.23	T10
	ATOM	11140	CA	SER	A	106	0.852	130.486	18.391	1.00	34.41	T10
5	ATOM	11141	CB	SER	A	106	1.496	129.401	19.256	1.00	23.86	T10
	ATOM	11142	OG	SER	A	106	2.875	129.248	18.954	1.00	25.97	T10
	ATOM	11143	C	SER	A	106	0.695	130.006	16.967	1.00	30.58	T10
	ATOM	11144	O	SER	A	106	-0.289	129.341	16.648	1.00	33.13	T10
	ATOM	11145	N	ALA	A	107	1.654	130.361	16.115	1.00	29.66	T10
10	ATOM	11146	CA	ALA	A	107	1.617	129.976	14.710	1.00	31.02	T10
	ATOM	11147	CB	ALA	A	107	0.734	130.944	13.930	1.00	29.22	T10
	ATOM	11148	C	ALA	A	107	3.014	129.940	14.105	1.00	31.05	T10
	ATOM	11149	O	ALA	A	107	3.963	130.462	14.676	1.00	32.72	T10
	ATOM	11150	N	GLY	A	108	3.132	129.317	12.943	1.00	37.60	T10
15	ATOM	11151	CA	GLY	A	108	4.416	129.231	12.282	1.00	34.41	T10
	ATOM	11152	C	GLY	A	108	4.273	128.642	10.892	1.00	29.74	T10
	ATOM	11153	O	GLY	A	108	3.166	128.310	10.478	1.00	35.72	T10
	ATOM	11154	N	ILE	A	109	5.383	128.514	10.170	1.00	29.81	T10
	ATOM	11155	CA	ILE	A	109	5.363	127.964	8.820	1.00	34.40	T10
20	ATOM	11156	CB	ILE	A	109	6.080	128.893	7.827	1.00	25.21	T10
	ATOM	11157	CG2	ILE	A	109	6.079	128.271	6.439	1.00	25.52	T10
	ATOM	11158	CG1	ILE	A	109	5.392	130.256	7.801	1.00	30.57	T10
	ATOM	11159	CD1	ILE	A	109	6.135	131.292	7.009	1.00	37.39	T10
	ATOM	11160	C	ILE	A	109	6.081	126.632	8.814	1.00	34.70	T10
25	ATOM	11161	O	ILE	A	109	7.097	126.479	9.481	1.00	24.52	T10
	ATOM	11162	N	ALA	A	110	5.554	125.672	8.062	1.00	31.45	T10
	ATOM	11163	CA	ALA	A	110	6.166	124.353	7.981	1.00	23.14	T10
	ATOM	11164	CB	ALA	A	110	5.587	123.442	9.032	1.00	29.29	T10
	ATOM	11165	C	ALA	A	110	5.916	123.767	6.614	1.00	25.81	T10
30	ATOM	11166	O	ALA	A	110	4.963	124.147	5.946	1.00	29.02	T10
	ATOM	11167	N	LYS	A	111	6.779	122.856	6.183	1.00	32.29	T10
	ATOM	11168	CA	LYS	A	111	6.582	122.231	4.891	1.00	26.31	T10
	ATOM	11169	CB	LYS	A	111	7.912	121.957	4.196	1.00	26.17	T10
	ATOM	11170	CG	LYS	A	111	7.718	121.348	2.816	1.00	33.29	T10
35	ATOM	11171	CD	LYS	A	111	9.038	121.122	2.099	1.00	32.47	T10
	ATOM	11172	CE	LYS	A	111	8.811	120.549	0.695	1.00	28.29	T10
	ATOM	11173	NZ	LYS	A	111	10.105	120.356	-0.058	1.00	36.34	T10
	ATOM	11174	C	LYS	A	111	5.850	120.924	5.148	1.00	26.80	T10
	ATOM	11175	O	LYS	A	111	6.229	120.176	6.045	1.00	29.68	T10
40	ATOM	11176	N	LEU	A	112	4.800	120.666	4.368	1.00	32.82	T10
	ATOM	11177	CA	LEU	A	112	3.992	119.461	4.520	1.00	34.81	T10
	ATOM	11178	CB	LEU	A	112	2.610	119.835	5.046	1.00	32.17	T10
	ATOM	11179	CG	LEU	A	112	2.569	120.749	6.271	1.00	27.75	T10
	ATOM	11180	CD1	LEU	A	112	1.161	121.252	6.484	1.00	34.14	T10
45	ATOM	11181	CD2	LEU	A	112	3.065	120.005	7.489	1.00	22.54	T10
	ATOM	11182	C	LEU	A	112	3.831	118.744	3.193	1.00	27.76	T10
	ATOM	11183	O	LEU	A	112	4.036	119.341	2.142	1.00	34.53	T10
	ATOM	11184	N	GLU	A	113	3.451	117.468	3.244	1.00	33.42	T10
	ATOM	11185	CA	GLU	A	113	3.248	116.676	2.034	1.00	27.92	T10
50	ATOM	11186	CB	GLU	A	113	4.282	115.566	1.915	1.00	27.15	T10
	ATOM	11187	CG	GLU	A	113	5.677	115.908	2.319	1.00	27.97	T10
	ATOM	11188	CD	GLU	A	113	6.682	114.931	1.729	1.00	22.73	T10
	ATOM	11189	OE1	GLU	A	113	6.430	113.696	1.777	1.00	31.88	T10
	ATOM	11190	OE2	GLU	A	113	7.726	115.402	1.208	1.00	32.76	T10
55	ATOM	11191	C	GLU	A	113	1.890	115.995	1.998	1.00	30.00	T10
	ATOM	11192	O	GLU	A	113	1.268	115.770	3.034	1.00	35.68	T10
	ATOM	11193	N	GLU	A	114	1.452	115.647	0.793	1.00	27.63	T10
	ATOM	11194	CA	GLU	A	114	0.195	114.936	0.605	1.00	39.45	T10
	ATOM	11195	CB	GLU	A	114	0.148	114.262	-0.752	1.00	30.98	T10
60	ATOM	11196	CG	GLU	A	114	-0.386	115.072	-1.883	1.00	28.37	T10
	ATOM	11197	CD	GLU	A	114	-0.813	114.168	-3.026	1.00	33.03	T10
	ATOM	11198	OE1	GLU	A	114	-1.753	113.354	-2.817	1.00	27.22	T10
	ATOM	11199	OE2	GLU	A	114	-0.202	114.265	-4.119	1.00	32.72	T10
	ATOM	11200	C	GLU	A	114	0.083	113.817	1.617	1.00	29.07	T10
65	ATOM	11201	O	GLU	A	114	0.958	112.962	1.688	1.00	28.78	T10
	ATOM	11202	N	GLY	A	115	-0.994	113.799	2.382	1.00	25.91	T10

	ATOM	11203	CA	GLY	A	115	-1.162	112.732	3.341	1.00	29.02	T10
	ATOM	11204	C	GLY	A	115	-0.794	113.097	4.755	1.00	30.45	T10
	ATOM	11205	O	GLY	A	115	-1.180	112.396	5.688	1.00	30.42	T10
	ATOM	11206	N	ASP	A	116	-0.025	114.164	4.936	1.00	32.73	T10
5	ATOM	11207	CA	ASP	A	116	0.326	114.551	6.294	1.00	38.09	T10
	ATOM	11208	CB	ASP	A	116	1.338	115.705	6.313	1.00	29.92	T10
	ATOM	11209	CG	ASP	A	116	2.722	115.292	5.854	1.00	33.99	T10
	ATOM	11210	OD1	ASP	A	116	3.074	114.104	6.020	1.00	29.65	T10
	ATOM	11211	OD2	ASP	A	116	3.469	116.165	5.349	1.00	31.58	T10
10	ATOM	11212	C	ASP	A	116	-0.958	115.003	6.987	1.00	34.72	T10
	ATOM	11213	O	ASP	A	116	-1.912	115.442	6.340	1.00	32.00	T10
	ATOM	11214	N	GLU	A	117	-0.991	114.884	8.303	1.00	26.71	T10
	ATOM	11215	CA	GLU	A	117	-2.156	115.321	9.043	1.00	33.28	T10
	ATOM	11216	CB	GLU	A	117	-2.881	114.130	9.653	1.00	30.40	T10
15	ATOM	11217	CG	GLU	A	117	-3.266	113.064	8.654	1.00	35.59	T10
	ATOM	11218	CD	GLU	A	117	-4.078	111.938	9.291	1.00	26.20	T10
	ATOM	11219	OE1	GLU	A	117	-3.792	111.579	10.471	1.00	31.43	T10
	ATOM	11220	OE2	GLU	A	117	-4.988	111.409	8.602	1.00	32.62	T10
	ATOM	11221	C	GLU	A	117	-1.699	116.255	10.150	1.00	30.57	T10
20	ATOM	11222	O	GLU	A	117	-0.654	116.034	10.763	1.00	37.57	T10
	ATOM	11223	N	LEU	A	118	-2.476	117.307	10.386	1.00	34.15	T10
	ATOM	11224	CA	LEU	A	118	-2.166	118.265	11.428	1.00	35.65	T10
	ATOM	11225	CB	LEU	A	118	-2.249	119.688	10.896	1.00	29.85	T10
	ATOM	11226	CG	LEU	A	118	-1.227	120.089	9.845	1.00	38.09	T10
25	ATOM	11227	CD1	LEU	A	118	-1.419	121.548	9.505	1.00	34.93	T10
	ATOM	11228	CD2	LEU	A	118	0.169	119.855	10.370	1.00	30.72	T10
	ATOM	11229	C	LEU	A	118	-3.186	118.097	12.530	1.00	33.05	T10
	ATOM	11230	O	LEU	A	118	-4.347	117.782	12.260	1.00	28.16	T10
	ATOM	11231	N	GLN	A	119	-2.759	118.303	13.771	1.00	29.48	T10
30	ATOM	11232	CA	GLN	A	119	-3.665	118.183	14.906	1.00	35.29	T10
	ATOM	11233	CB	GLN	A	119	-3.735	116.732	15.366	1.00	32.78	T10
	ATOM	11234	CG	GLN	A	119	-2.423	116.184	15.875	1.00	31.23	T10
	ATOM	11235	CD	GLN	A	119	-2.518	114.718	16.251	1.00	29.55	T10
	ATOM	11236	OE1	GLN	A	119	-1.671	114.202	16.983	1.00	27.43	T10
35	ATOM	11237	NE2	GLN	A	119	-3.547	114.034	15.744	1.00	27.96	T10
	ATOM	11238	C	GLN	A	119	-3.231	119.082	16.060	1.00	26.21	T10
	ATOM	11239	O	GLN	A	119	-2.056	119.426	16.182	1.00	32.30	T10
	ATOM	11240	N	LEU	A	120	-4.196	119.456	16.896	1.00	34.17	T10
	ATOM	11241	CA	LEU	A	120	-3.967	120.328	18.046	1.00	21.88	T10
40	ATOM	11242	CB	LEU	A	120	-5.002	121.463	18.022	1.00	32.05	T10
	ATOM	11243	CG	LEU	A	120	-4.954	122.725	18.893	1.00	33.12	T10
	ATOM	11244	CD1	LEU	A	120	-4.623	122.372	20.333	1.00	33.08	T10
	ATOM	11245	CD2	LEU	A	120	-3.934	123.677	18.330	1.00	35.99	T10
	ATOM	11246	C	LEU	A	120	-4.153	119.475	19.293	1.00	24.24	T10
45	ATOM	11247	O	LEU	A	120	-5.222	118.907	19.503	1.00	28.30	T10
	ATOM	11248	N	ALA	A	121	-3.129	119.391	20.132	1.00	33.20	T10
	ATOM	11249	CA	ALA	A	121	-3.234	118.561	21.327	1.00	34.22	T10
	ATOM	11250	CB	ALA	A	121	-2.396	117.296	21.140	1.00	34.75	T10
	ATOM	11251	C	ALA	A	121	-2.852	119.237	22.641	1.00	40.50	T10
50	ATOM	11252	O	ALA	A	121	-1.905	120.021	22.708	1.00	32.27	T10
	ATOM	11253	N	ILE	A	122	-3.603	118.913	23.688	1.00	29.67	T10
	ATOM	11254	CA	ILE	A	122	-3.345	119.449	25.016	1.00	26.07	T10
	ATOM	11255	CB	ILE	A	122	-4.625	120.001	25.670	1.00	31.39	T10
	ATOM	11256	CG2	ILE	A	122	-4.286	120.582	27.034	1.00	29.28	T10
55	ATOM	11257	CG1	ILE	A	122	-5.246	121.074	24.767	1.00	20.57	T10
	ATOM	11258	CD1	ILE	A	122	-6.501	121.697	25.312	1.00	28.35	T10
	ATOM	11259	C	ILE	A	122	-2.789	118.320	25.880	1.00	31.14	T10
	ATOM	11260	O	ILE	A	122	-3.450	117.302	26.105	1.00	26.24	T10
	ATOM	11261	N	PRO	A	123	-1.554	118.487	26.369	1.00	40.34	T10
60	ATOM	11262	CD	PRO	A	123	-0.662	119.608	26.043	1.00	26.25	T10
	ATOM	11263	CA	PRO	A	123	-0.859	117.506	27.213	1.00	28.15	T10
	ATOM	11264	CB	PRO	A	123	0.587	118.008	27.220	1.00	33.65	T10
	ATOM	11265	CG	PRO	A	123	0.666	118.922	26.020	1.00	29.31	T10
	ATOM	11266	C	PRO	A	123	-1.416	117.409	28.635	1.00	33.52	T10
65	ATOM	11267	O	PRO	A	123	-0.682	117.585	29.610	1.00	38.38	T10
	ATOM	11268	N	ARG	A	124	-2.712	117.149	28.756	1.00	29.69	T10

	ATOM	11269	CA	ARG A 124	-3.341	117.012	30.064	1.00	31.89	T10
	ATOM	11270	CB	ARG A 124	-3.875	118.336	30.569	1.00	22.15	T10
	ATOM	11271	CG	ARG A 124	-2.841	119.149	31.294	1.00	31.91	T10
	ATOM	11272	CD	ARG A 124	-3.467	119.742	32.537	1.00	34.31	T10
5	ATOM	11273	NE	ARG A 124	-3.693	118.736	33.578	1.00	30.97	T10
	ATOM	11274	CZ	ARG A 124	-4.392	118.966	34.689	1.00	29.97	T10
	ATOM	11275	NH1	ARG A 124	-4.934	120.166	34.889	1.00	34.69	T10
	ATOM	11276	NH2	ARG A 124	-4.540	118.009	35.605	1.00	32.81	T10
	ATOM	11277	C	ARG A 124	-4.476	116.040	29.948	1.00	31.58	T10
10	ATOM	11278	O	ARG A 124	-5.015	115.843	28.856	1.00	33.79	T10
	ATOM	11279	N	GLU A 125	-4.854	115.442	31.071	1.00	31.34	T10
	ATOM	11280	CA	GLU A 125	-5.920	114.455	31.040	1.00	31.04	T10
	ATOM	11281	CB	GLU A 125	-5.875	113.591	32.292	1.00	27.57	T10
	ATOM	11282	CG	GLU A 125	-4.691	112.626	32.283	1.00	40.77	T10
15	ATOM	11283	CD	GLU A 125	-5.068	111.238	32.798	1.00	41.01	T10
	ATOM	11284	OE1	GLU A 125	-5.584	111.155	33.945	1.00	34.76	T10
	ATOM	11285	OE2	GLU A 125	-4.848	110.237	32.056	1.00	28.40	T10
	ATOM	11286	C	GLU A 125	-7.319	115.009	30.819	1.00	30.86	T10
	ATOM	11287	O	GLU A 125	-8.048	114.517	29.952	1.00	32.08	T10
20	ATOM	11288	N	ASN A 126	-7.717	116.012	31.587	1.00	29.60	T10
	ATOM	11289	CA	ASN A 126	-9.036	116.581	31.358	1.00	31.43	T10
	ATOM	11290	CB	ASN A 126	-10.054	116.066	32.374	1.00	20.24	T10
	ATOM	11291	CG	ASN A 126	-10.604	114.694	31.998	1.00	24.47	T10
	ATOM	11292	OD1	ASN A 126	-9.954	113.659	32.214	1.00	27.28	T10
25	ATOM	11293	ND2	ASN A 126	-11.806	114.681	31.411	1.00	34.52	T10
	ATOM	11294	C	ASN A 126	-8.929	118.079	31.433	1.00	26.94	T10
	ATOM	11295	O	ASN A 126	-9.525	118.725	32.300	1.00	32.46	T10
	ATOM	11296	N	ALA A 127	-8.150	118.623	30.505	1.00	29.13	T10
	ATOM	11297	CA	ALA A 127	-7.910	120.052	30.438	1.00	33.40	T10
30	ATOM	11298	CB	ALA A 127	-7.207	120.384	29.133	1.00	34.61	T10
	ATOM	11299	C	ALA A 127	-9.199	120.856	30.559	1.00	31.73	T10
	ATOM	11300	O	ALA A 127	-10.193	120.551	29.894	1.00	29.14	T10
	ATOM	11301	N	GLN A 128	-9.183	121.862	31.431	1.00	30.24	T10
	ATOM	11302	CA	GLN A 128	-10.338	122.742	31.601	1.00	30.45	T10
35	ATOM	11303	CB	GLN A 128	-10.328	123.376	32.993	1.00	28.40	T10
	ATOM	11304	CG	GLN A 128	-10.536	122.347	34.098	1.00	30.98	T10
	ATOM	11305	CD	GLN A 128	-11.686	121.392	33.773	1.00	39.04	T10
	ATOM	11306	OE1	GLN A 128	-12.831	121.812	33.569	1.00	27.03	T10
	ATOM	11307	NE2	GLN A 128	-11.381	120.103	33.714	1.00	32.14	T10
40	ATOM	11308	C	GLN A 128	-10.200	123.802	30.516	1.00	34.23	T10
	ATOM	11309	O	GLN A 128	-9.443	124.768	30.652	1.00	38.27	T10
	ATOM	11310	N	ILE A 129	-10.946	123.601	29.440	1.00	29.62	T10
	ATOM	11311	CA	ILE A 129	-10.893	124.455	28.264	1.00	27.26	T10
	ATOM	11312	CB	ILE A 129	-10.384	123.570	27.072	1.00	30.59	T10
45	ATOM	11313	CG2	ILE A 129	-11.285	123.665	25.861	1.00	28.08	T10
	ATOM	11314	CG1	ILE A 129	-8.937	123.919	26.771	1.00	28.02	T10
	ATOM	11315	CD1	ILE A 129	-8.031	123.672	27.938	1.00	35.87	T10
	ATOM	11316	C	ILE A 129	-12.224	125.126	27.916	1.00	24.24	T10
	ATOM	11317	O	ILE A 129	-13.282	124.676	28.343	1.00	31.09	T10
50	ATOM	11318	N	SER A 130	-12.171	126.213	27.160	1.00	30.88	T10
	ATOM	11319	CA	SER A 130	-13.392	126.874	26.726	1.00	29.45	T10
	ATOM	11320	CB	SER A 130	-13.244	128.387	26.796	1.00	37.65	T10
	ATOM	11321	OG	SER A 130	-14.331	129.013	26.135	1.00	28.50	T10
	ATOM	11322	C	SER A 130	-13.625	126.459	25.278	1.00	34.14	T10
55	ATOM	11323	O	SER A 130	-12.737	126.604	24.440	1.00	28.86	T10
	ATOM	11324	N	LEU A 131	-14.805	125.941	24.968	1.00	28.83	T10
	ATOM	11325	CA	LEU A 131	-15.082	125.522	23.594	1.00	31.94	T10
	ATOM	11326	CB	LEU A 131	-15.836	124.194	23.575	1.00	28.23	T10
	ATOM	11327	CG	LEU A 131	-14.986	122.949	23.812	1.00	30.21	T10
60	ATOM	11328	CD1	LEU A 131	-14.285	123.039	25.144	1.00	30.26	T10
	ATOM	11329	CD2	LEU A 131	-15.882	121.730	23.767	1.00	34.47	T10
	ATOM	11330	C	LEU A 131	-15.843	126.540	22.748	1.00	28.97	T10
	ATOM	11331	O	LEU A 131	-16.633	126.166	21.879	1.00	34.99	T10
	ATOM	11332	N	ASP A 132	-15.595	127.823	22.994	1.00	26.61	T10
65	ATOM	11333	CA	ASP A 132	-16.253	128.874	22.229	1.00	32.27	T10
	ATOM	11334	CB	ASP A 132	-16.377	130.146	23.068	1.00	34.10	T10

	ATOM	11335	CG	ASP	A	132	-17.590	130.117	23.997	1.00	28.69	T10
	ATOM	11336	OD1	ASP	A	132	-17.673	131.004	24.887	1.00	30.34	T10
	ATOM	11337	OD2	ASP	A	132	-18.459	129.217	23.825	1.00	29.53	T10
	ATOM	11338	C	ASP	A	132	-15.525	129.185	20.927	1.00	30.50	T10
5	ATOM	11339	O	ASP	A	132	-14.297	129.196	20.866	1.00	38.19	T10
	ATOM	11340	N	GLY	A	133	-16.308	129.437	19.886	1.00	31.38	T10
	ATOM	11341	CA	GLY	A	133	-15.757	129.749	18.583	1.00	33.56	T10
	ATOM	11342	C	GLY	A	133	-14.747	130.878	18.551	1.00	33.04	T10
	ATOM	11343	O	GLY	A	133	-13.840	130.842	17.725	1.00	39.84	T10
10	ATOM	11344	N	ASP	A	134	-14.887	131.876	19.424	1.00	29.80	T10
	ATOM	11345	CA	ASP	A	134	-13.945	132.997	19.441	1.00	35.17	T10
	ATOM	11346	CB	ASP	A	134	-14.371	134.124	20.372	1.00	27.47	T10
	ATOM	11347	CG	ASP	A	134	-15.823	134.178	20.592	1.00	34.37	T10
	ATOM	11348	OD1	ASP	A	134	-16.533	134.408	19.594	1.00	28.43	T10
15	ATOM	11349	OD2	ASP	A	134	-16.239	133.997	21.760	1.00	27.18	T10
	ATOM	11350	C	ASP	A	134	-12.603	132.584	19.983	1.00	28.65	T10
	ATOM	11351	O	ASP	A	134	-11.575	132.729	19.338	1.00	38.24	T10
	ATOM	11352	N	VAL	A	135	-12.633	132.099	21.210	1.00	35.84	T10
	ATOM	11353	CA	VAL	A	135	-11.431	131.732	21.915	1.00	31.47	T10
20	ATOM	11354	CB	VAL	A	135	-11.768	131.564	23.385	1.00	24.02	T10
	ATOM	11355	CG1	VAL	A	135	-12.314	132.882	23.915	1.00	34.36	T10
	ATOM	11356	CG2	VAL	A	135	-12.803	130.453	23.554	1.00	22.80	T10
	ATOM	11357	C	VAL	A	135	-10.568	130.564	21.450	1.00	29.24	T10
	ATOM	11358	O	VAL	A	135	-9.351	130.655	21.544	1.00	35.77	T10
25	ATOM	11359	N	THR	A	136	-11.146	129.469	20.966	1.00	27.64	T10
	ATOM	11360	CA	THR	A	136	-10.284	128.372	20.529	1.00	33.16	T10
	ATOM	11361	CB	THR	A	136	-10.357	127.168	21.515	1.00	25.69	T10
	ATOM	11362	OG1	THR	A	136	-11.557	126.434	21.297	1.00	38.40	T10
	ATOM	11363	CG2	THR	A	136	-10.361	127.652	22.942	1.00	28.40	T10
30	ATOM	11364	C	THR	A	136	-10.553	127.916	19.087	1.00	27.88	T10
	ATOM	11365	O	THR	A	136	-11.663	127.503	18.739	1.00	33.07	T10
	ATOM	11366	N	PHE	A	137	-9.518	128.008	18.256	1.00	27.05	T10
	ATOM	11367	CA	PHE	A	137	-9.616	127.645	16.853	1.00	34.70	T10
	ATOM	11368	CB	PHE	A	137	-10.024	128.878	16.049	1.00	34.66	T10
35	ATOM	11369	CG	PHE	A	137	-9.285	130.132	16.434	1.00	27.31	T10
	ATOM	11370	CD1	PHE	A	137	-8.006	130.385	15.956	1.00	24.01	T10
	ATOM	11371	CD2	PHE	A	137	-9.868	131.063	17.272	1.00	28.17	T10
	ATOM	11372	CE1	PHE	A	137	-7.327	131.547	16.308	1.00	34.42	T10
	ATOM	11373	CE2	PHE	A	137	-9.194	132.220	17.624	1.00	33.30	T10
40	ATOM	11374	CZ	PHE	A	137	-7.926	132.460	17.142	1.00	35.58	T10
	ATOM	11375	C	PHE	A	137	-8.319	127.041	16.309	1.00	27.13	T10
	ATOM	11376	O	PHE	A	137	-7.267	127.162	16.927	1.00	23.42	T10
	ATOM	11377	N	PHE	A	138	-8.396	126.403	15.144	1.00	29.09	T10
	ATOM	11378	CA	PHE	A	138	-7.239	125.745	14.546	1.00	22.72	T10
45	ATOM	11379	CB	PHE	A	138	-7.442	124.227	14.668	1.00	32.29	T10
	ATOM	11380	CG	PHE	A	138	-6.213	123.417	14.414	1.00	33.05	T10
	ATOM	11381	CD1	PHE	A	138	-4.944	123.970	14.560	1.00	35.43	T10
	ATOM	11382	CD2	PHE	A	138	-6.324	122.093	14.023	1.00	32.07	T10
	ATOM	11383	CE1	PHE	A	138	-3.806	123.210	14.311	1.00	33.09	T10
50	ATOM	11384	CE2	PHE	A	138	-5.191	121.324	13.773	1.00	33.06	T10
	ATOM	11385	CZ	PHE	A	138	-3.932	121.880	13.915	1.00	30.40	T10
	ATOM	11386	C	PHE	A	138	-6.991	126.187	13.093	1.00	29.87	T10
	ATOM	11387	O	PHE	A	138	-7.895	126.178	12.262	1.00	33.58	T10
	ATOM	11388	N	GLY	A	139	-5.732	126.540	12.823	1.00	30.17	T10
55	ATOM	11389	CA	GLY	A	139	-5.279	127.080	11.541	1.00	26.27	T10
	ATOM	11390	C	GLY	A	139	-5.102	126.352	10.220	1.00	31.39	T10
	ATOM	11391	O	GLY	A	139	-5.990	125.629	9.793	1.00	29.26	T10
	ATOM	11392	N	ALA	A	140	-3.974	126.609	9.552	1.00	31.01	T10
	ATOM	11393	CA	ALA	A	140	-3.609	126.016	8.252	1.00	37.52	T10
60	ATOM	11394	CB	ALA	A	140	-4.095	124.589	8.163	1.00	30.51	T10
	ATOM	11395	C	ALA	A	140	-4.037	126.771	6.989	1.00	34.37	T10
	ATOM	11396	O	ALA	A	140	-5.195	126.745	6.599	1.00	27.61	T10
	ATOM	11397	N	LEU	A	141	-3.067	127.419	6.347	1.00	26.69	T10
	ATOM	11398	CA	LEU	A	141	-3.264	128.187	5.116	1.00	32.38	T10
65	ATOM	11399	CB	LEU	A	141	-3.365	129.687	5.433	1.00	29.32	T10
	ATOM	11400	CG	LEU	A	141	-3.397	130.704	4.284	1.00	35.61	T10

TABLE 3

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5	ATOM	1	CB	VAL	1	-10.775	93.262	-41.416	1.00 69.28
	ATOM	2	CG1	VAL	1	-11.154	94.190	-40.241	1.00 68.26
	ATOM	3	CG2	VAL	1	-10.256	94.066	-42.613	1.00 71.75
	ATOM	4	C	VAL	1	-10.392	91.191	-40.047	1.00 64.86
	ATOM	5	O	VAL	1	-10.228	91.222	-38.818	1.00 64.02
10	ATOM	6	N	VAL	1	-9.057	91.553	-42.128	1.00 65.10
	ATOM	7	CA	VAL	1	-9.700	92.224	-40.958	1.00 65.98
	ATOM	8	N	THR	2	-11.172	90.289	-40.648	1.00 61.42
	ATOM	9	CA	THR	2	-11.875	89.260	-39.883	1.00 56.64
	ATOM	10	CB	THR	2	-13.401	89.352	-40.059	1.00 56.77
15	ATOM	11	OG1	THR	2	-13.714	89.295	-41.453	1.00 57.45
	ATOM	12	CG2	THR	2	-13.943	90.648	-39.467	1.00 55.68
	ATOM	13	C	THR	2	-11.463	87.865	-40.312	1.00 53.08
	ATOM	14	O	THR	2	-10.794	87.684	-41.324	1.00 52.00
	ATOM	15	N	GLN	3	-11.888	86.875	-39.538	1.00 50.34
20	ATOM	16	CA	GLN	3	-11.568	85.484	-39.833	1.00 45.97
	ATOM	17	CB	GLN	3	-10.949	84.819	-38.608	1.00 46.09
	ATOM	18	CG	GLN	3	-9.837	85.615	-37.978	1.00 49.60
	ATOM	19	CD	GLN	3	-9.204	84.882	-36.823	1.00 50.43
	ATOM	20	OE1	GLN	3	-8.553	83.861	-37.016	1.00 50.94
25	ATOM	21	NE2	GLN	3	-9.397	85.395	-35.609	1.00 49.99
	ATOM	22	C	GLN	3	-12.811	84.717	-40.242	1.00 42.41
	ATOM	23	O	GLN	3	-13.690	84.450	-39.428	1.00 37.34
	ATOM	24	N	ASP	4	-12.882	84.362	-41.515	1.00 40.32
	ATOM	25	CA	ASP	4	-14.024	83.621	-42.018	1.00 39.37
30	ATOM	26	CB	ASP	4	-13.963	83.532	-43.546	1.00 44.56
	ATOM	27	CG	ASP	4	-14.109	84.884	-44.213	1.00 45.76
	ATOM	28	OD1	ASP	4	-13.893	85.907	-43.523	1.00 48.58
	ATOM	29	OD2	ASP	4	-14.429	84.923	-45.420	1.00 45.55
	ATOM	30	C	ASP	4	-14.009	82.228	-41.429	1.00 36.30
35	ATOM	31	O	ASP	4	-12.949	81.660	-41.178	1.00 36.29
	ATOM	32	N	CYS	5	-15.193	81.683	-41.197	1.00 35.11
	ATOM	33	CA	CYS	5	-15.314	80.337	-40.667	1.00 35.61
	ATOM	34	CB	CYS	5	-15.011	80.304	-39.167	1.00 34.96
	ATOM	35	SG	CYS	5	-15.711	81.640	-38.218	1.00 35.34
40	ATOM	36	C	CYS	5	-16.702	79.803	-40.949	1.00 31.80
	ATOM	37	O	CYS	5	-17.657	80.561	-41.069	1.00 31.85
	ATOM	38	N	LEU	6	-16.796	78.491	-41.094	1.00 27.72
	ATOM	39	CA	LEU	6	-18.059	77.833	-41.357	1.00 30.80
	ATOM	40	CB	LEU	6	-18.177	77.489	-42.837	1.00 29.25
45	ATOM	41	CG	LEU	6	-19.419	76.719	-43.291	1.00 29.47
	ATOM	42	CD1	LEU	6	-19.682	76.995	-44.747	1.00 30.65
	ATOM	43	CD2	LEU	6	-19.211	75.249	-43.081	1.00 35.27
	ATOM	44	C	LEU	6	-18.090	76.573	-40.515	1.00 31.83
	ATOM	45	O	LEU	6	-17.158	75.779	-40.558	1.00 33.45
50	ATOM	46	N	GLN	7	-19.153	76.393	-39.740	1.00 32.23
	ATOM	47	CA	GLN	7	-19.265	75.228	-38.889	1.00 30.26
	ATOM	48	CB	GLN	7	-19.343	75.664	-37.432	1.00 29.56
	ATOM	49	CG	GLN	7	-19.167	74.551	-36.422	1.00 25.09
	ATOM	50	CD	GLN	7	-18.999	75.091	-35.016	1.00 26.50
55	ATOM	51	OE1	GLN	7	-19.942	75.603	-34.412	1.00 24.40
	ATOM	52	NE2	GLN	7	-17.790	74.994	-34.495	1.00 27.03
	ATOM	53	C	GLN	7	-20.486	74.418	-39.266	1.00 32.11
	ATOM	54	O	GLN	7	-21.534	74.966	-39.584	1.00 34.24
	ATOM	55	N	LEU	8	-20.327	73.103	-39.241	1.00 33.73
60	ATOM	56	CA	LEU	8	-21.398	72.178	-39.573	1.00 35.41
	ATOM	57	CB	LEU	8	-20.962	71.264	-40.732	1.00 32.44
	ATOM	58	CG	LEU	8	-21.286	71.633	-42.187	1.00 27.69
	ATOM	59	CD1	LEU	8	-21.617	73.096	-42.332	1.00 26.43
	ATOM	60	CD2	LEU	8	-20.114	71.244	-43.047	1.00 18.64
65	ATOM	61	C	LEU	8	-21.766	71.345	-38.354	1.00 36.87
	ATOM	62	O	LEU	8	-20.923	71.025	-37.518	1.00 36.05
	ATOM	63	N	ILE	9	-23.040	70.991	-38.273	1.00 39.25

	ATOM	64	CA	ILE	9	-23.572	70.215	-37.165	1.00	35.57
	ATOM	65	CB	ILE	9	-24.533	71.113	-36.347	1.00	34.28
	ATOM	66	CG2	ILE	9	-25.821	70.401	-36.018	1.00	38.20
	ATOM	67	CG1	ILE	9	-23.839	71.581	-35.088	1.00	35.22
5	ATOM	68	CD1	ILE	9	-24.746	72.402	-34.235	1.00	43.63
	ATOM	69	C	ILE	9	-24.292	68.977	-37.707	1.00	35.28
	ATOM	70	O	ILE	9	-24.873	69.017	-38.788	1.00	33.75
	ATOM	71	N	ALA	10	-24.242	67.873	-36.971	1.00	35.78
	ATOM	72	CA	ALA	10	-24.914	66.663	-37.420	1.00	35.28
10	ATOM	73	CB	ALA	10	-24.620	65.519	-36.477	1.00	28.47
	ATOM	74	C	ALA	10	-26.418	66.889	-37.506	1.00	36.49
	ATOM	75	O	ALA	10	-27.014	67.531	-36.646	1.00	37.41
	ATOM	76	N	ASP	11	-27.028	66.359	-38.556	1.00	39.44
	ATOM	77	CA	ASP	11	-28.466	66.493	-38.755	1.00	40.77
15	ATOM	78	CB	ASP	11	-28.758	66.896	-40.202	1.00	38.90
	ATOM	79	CG	ASP	11	-30.237	66.923	-40.504	1.00	40.34
	ATOM	80	OD1	ASP	11	-31.041	67.118	-39.562	1.00	39.50
	ATOM	81	OD2	ASP	11	-30.589	66.760	-41.688	1.00	37.38
	ATOM	82	C	ASP	11	-29.165	65.179	-38.422	1.00	42.25
20	ATOM	83	O	ASP	11	-29.287	64.288	-39.263	1.00	40.54
	ATOM	84	N	SER	12	-29.628	65.078	-37.182	1.00	42.92
	ATOM	85	CA	SER	12	-30.288	63.874	-36.683	1.00	43.84
	ATOM	86	CB	SER	12	-30.569	64.026	-35.193	1.00	42.13
	ATOM	87	OG	SER	12	-31.375	65.167	-34.955	1.00	43.31
25	ATOM	88	C	SER	12	-31.581	63.503	-37.390	1.00	46.10
	ATOM	89	O	SER	12	-32.169	62.459	-37.109	1.00	43.90
	ATOM	90	N	GLU	13	-32.020	64.352	-38.313	1.00	46.72
	ATOM	91	CA	GLU	13	-33.254	64.087	-39.027	1.00	44.93
	ATOM	92	CB	GLU	13	-34.102	65.346	-39.072	1.00	46.71
30	ATOM	93	CG	GLU	13	-34.592	65.736	-37.704	1.00	58.25
	ATOM	94	CD	GLU	13	-35.858	66.565	-37.765	1.00	66.70
	ATOM	95	OE1	GLU	13	-36.861	66.058	-38.341	1.00	74.10
	ATOM	96	OE2	GLU	13	-35.858	67.712	-37.243	1.00	71.55
	ATOM	97	C	GLU	13	-33.110	63.491	-40.423	1.00	43.24
35	ATOM	98	O	GLU	13	-34.073	63.459	-41.192	1.00	44.25
	ATOM	99	N	THR	14	-31.914	63.030	-40.760	1.00	37.06
	ATOM	100	CA	THR	14	-31.716	62.376	-42.047	1.00	36.77
	ATOM	101	CB	THR	14	-31.148	63.333	-43.137	1.00	33.39
	ATOM	102	OG1	THR	14	-29.765	63.583	-42.896	1.00	41.69
40	ATOM	103	CG2	THR	14	-31.906	64.651	-43.140	1.00	31.66
	ATOM	104	C	THR	14	-30.753	61.220	-41.798	1.00	34.96
	ATOM	105	O	THR	14	-29.919	61.279	-40.902	1.00	35.04
	ATOM	106	N	PRO	15	-30.873	60.145	-42.577	1.00	37.11
	ATOM	107	CD	PRO	15	-31.855	59.961	-43.655	1.00	36.50
45	ATOM	108	CA	PRO	15	-30.019	58.960	-42.439	1.00	38.81
	ATOM	109	CB	PRO	15	-30.619	57.982	-43.449	1.00	39.15
	ATOM	110	CG	PRO	15	-32.032	58.478	-43.641	1.00	37.32
	ATOM	111	C	PRO	15	-28.555	59.244	-42.751	1.00	37.78
	ATOM	112	O	PRO	15	-28.252	60.137	-43.536	1.00	35.91
50	ATOM	113	N	THR	16	-27.646	58.487	-42.145	1.00	36.67
	ATOM	114	CA	THR	16	-26.235	58.694	-42.428	1.00	38.46
	ATOM	115	CB	THR	16	-25.328	57.855	-41.507	1.00	38.61
	ATOM	116	OG1	THR	16	-25.558	56.462	-41.744	1.00	42.44
	ATOM	117	CG2	THR	16	-25.618	58.172	-40.048	1.00	40.11
55	ATOM	118	C	THR	16	-26.030	58.235	-43.855	1.00	35.97
	ATOM	119	O	THR	16	-26.577	57.216	-44.254	1.00	39.85
	ATOM	120	N	ILE	17	-25.252	58.985	-44.625	1.00	33.66
	ATOM	121	CA	ILE	17	-25.005	58.632	-46.016	1.00	34.76
	ATOM	122	CB	ILE	17	-24.467	59.840	-46.793	1.00	32.44
60	ATOM	123	CG2	ILE	17	-24.099	59.430	-48.206	1.00	28.66
	ATOM	124	CG1	ILE	17	-25.516	60.957	-46.791	1.00	33.15
	ATOM	125	CD1	ILE	17	-25.069	62.240	-47.464	1.00	31.19
	ATOM	126	C	ILE	17	-24.035	57.471	-46.184	1.00	40.64
	ATOM	127	O	ILE	17	-22.939	57.483	-45.639	1.00	44.07
65	ATOM	128	N	GLN	18	-24.445	56.459	-46.945	1.00	45.91
	ATOM	129	CA	GLN	18	-23.600	55.294	-47.194	1.00	50.22

	ATOM	130	CB	GLN	18	-24.397	54.004	-47.033	1.00	53.09
	ATOM	131	CG	GLN	18	-23.798	53.074	-46.011	1.00	57.60
	ATOM	132	CD	GLN	18	-23.923	53.644	-44.622	1.00	61.39
	ATOM	133	OE1	GLN	18	-25.025	53.738	-44.076	1.00	68.36
5	ATOM	134	NE2	GLN	18	-22.800	54.048	-44.045	1.00	64.93
	ATOM	135	C	GLN	18	-23.030	55.338	-48.600	1.00	51.56
	ATOM	136	O	GLN	18	-23.751	55.573	-49.565	1.00	56.44
	ATOM	137	N	LYS	19	-21.734	55.102	-48.720	1.00	52.60
	ATOM	138	CA	LYS	19	-21.109	55.122	-50.032	1.00	56.10
10	ATOM	139	CB	LYS	19	-21.073	56.546	-50.588	1.00	55.55
	ATOM	140	CG	LYS	19	-20.331	56.619	-51.902	1.00	61.15
	ATOM	141	CD	LYS	19	-20.464	57.962	-52.588	1.00	64.42
	ATOM	142	CE	LYS	19	-19.701	57.952	-53.922	1.00	64.84
	ATOM	143	NZ	LYS	19	-19.822	59.244	-54.670	1.00	69.52
15	ATOM	144	C	LYS	19	-19.699	54.551	-50.018	1.00	57.41
	ATOM	145	O	LYS	19	-18.899	54.878	-49.140	1.00	61.02
	ATOM	146	N	GLY	20	-19.404	53.704	-51.003	1.00	58.84
	ATOM	147	CA	GLY	20	-18.093	53.091	-51.107	1.00	56.66
	ATOM	148	C	GLY	20	-17.615	52.487	-49.800	1.00	56.87
20	ATOM	149	O	GLY	20	-16.427	52.568	-49.485	1.00	58.64
	ATOM	150	N	SER	21	-18.537	51.883	-49.049	1.00	54.49
	ATOM	151	CA	SER	21	-18.225	51.257	-47.761	1.00	53.68
	ATOM	152	CB	SER	21	-17.213	50.119	-47.949	1.00	55.80
	ATOM	153	OG	SER	21	-15.880	50.595	-47.957	1.00	65.34
25	ATOM	154	C	SER	21	-17.703	52.273	-46.716	1.00	51.41
	ATOM	155	O	SER	21	-16.952	51.926	-45.789	1.00	47.74
	ATOM	156	N	TYR	22	-18.111	53.530	-46.889	1.00	46.25
	ATOM	157	CA	TYR	22	-17.751	54.615	-45.987	1.00	40.57
	ATOM	158	CB	TYR	22	-16.962	55.696	-46.731	1.00	43.27
30	ATOM	159	CG	TYR	22	-15.475	55.456	-46.774	1.00	49.18
	ATOM	160	CD1	TYR	22	-14.950	54.307	-47.356	1.00	52.61
	ATOM	161	CE1	TYR	22	-13.572	54.075	-47.386	1.00	51.92
	ATOM	162	CD2	TYR	22	-14.587	56.373	-46.223	1.00	50.01
	ATOM	163	CE2	TYR	22	-13.209	56.152	-46.251	1.00	50.94
35	ATOM	164	CZ	TYR	22	-12.710	55.002	-46.833	1.00	50.78
	ATOM	165	OH	TYR	22	-11.351	54.782	-46.857	1.00	53.87
	ATOM	166	C	TYR	22	-19.044	55.220	-45.452	1.00	37.85
	ATOM	167	O	TYR	22	-20.065	55.213	-46.139	1.00	38.70
	ATOM	168	N	THR	23	-19.015	55.725	-44.226	1.00	33.81
40	ATOM	169	CA	THR	23	-20.202	56.351	-43.655	1.00	32.40
	ATOM	170	CB	THR	23	-20.505	55.860	-42.216	1.00	31.58
	ATOM	171	OG1	THR	23	-20.650	54.437	-42.202	1.00	30.76
	ATOM	172	CG2	THR	23	-21.783	56.493	-41.709	1.00	24.08
	ATOM	173	C	THR	23	-19.951	57.847	-43.594	1.00	33.28
45	ATOM	174	O	THR	23	-18.902	58.283	-43.132	1.00	30.44
	ATOM	175	N	PHE	24	-20.912	58.628	-44.063	1.00	32.16
	ATOM	176	CA	PHE	24	-20.784	60.075	-44.044	1.00	30.68
	ATOM	177	CB	PHE	24	-20.837	60.637	-45.465	1.00	28.20
	ATOM	178	CG	PHE	24	-19.672	60.236	-46.313	1.00	31.00
50	ATOM	179	CD1	PHE	24	-19.667	59.022	-46.979	1.00	31.69
	ATOM	180	CD2	PHE	24	-18.558	61.063	-46.423	1.00	31.42
	ATOM	181	CE1	PHE	24	-18.571	58.636	-47.740	1.00	32.57
	ATOM	182	CE2	PHE	24	-17.462	60.685	-47.180	1.00	25.01
	ATOM	183	CZ	PHE	24	-17.469	59.469	-47.839	1.00	30.12
55	ATOM	184	C	PHE	24	-21.867	60.726	-43.194	1.00	31.73
	ATOM	185	O	PHE	24	-23.050	60.438	-43.343	1.00	33.31
	ATOM	186	N	VAL	25	-21.449	61.602	-42.295	1.00	32.92
	ATOM	187	CA	VAL	25	-22.379	62.300	-41.434	1.00	30.84
	ATOM	188	CB	VAL	25	-21.634	63.083	-40.341	1.00	30.29
60	ATOM	189	CG1	VAL	25	-22.618	63.837	-39.481	1.00	28.13
	ATOM	190	CG2	VAL	25	-20.809	62.143	-39.505	1.00	29.53
	ATOM	191	C	VAL	25	-23.200	63.293	-42.248	1.00	36.08
	ATOM	192	O	VAL	25	-22.669	63.994	-43.106	1.00	35.82
	ATOM	193	N	PRO	26	-24.518	63.345	-42.007	1.00	38.03
65	ATOM	194	CD	PRO	26	-25.293	62.356	-41.242	1.00	38.28
	ATOM	195	CA	PRO	26	-25.418	64.266	-42.716	1.00	38.08

	ATOM	196	CB	PRO	26	-26.806	63.674	-42.456	1.00	37.62
	ATOM	197	CG	PRO	26	-26.539	62.257	-42.062	1.00	36.31
	ATOM	198	C	PRO	26	-25.262	65.620	-42.027	1.00	40.85
	ATOM	199	O	PRO	26	-25.664	65.778	-40.875	1.00	42.27
5	ATOM	200	N	TRP	27	-24.686	66.599	-42.709	1.00	40.86
	ATOM	201	CA	TRP	27	-24.479	67.891	-42.067	1.00	37.05
	ATOM	202	CB	TRP	27	-23.196	68.545	-42.585	1.00	33.29
	ATOM	203	CG	TRP	27	-21.974	67.739	-42.326	1.00	31.06
	ATOM	204	CD2	TRP	27	-21.457	67.350	-41.053	1.00	27.79
10	ATOM	205	CE2	TRP	27	-20.312	66.572	-41.291	1.00	29.28
	ATOM	206	CE3	TRP	27	-21.851	67.582	-39.735	1.00	27.06
	ATOM	207	CD1	TRP	27	-21.146	67.203	-43.255	1.00	31.07
	ATOM	208	NE1	TRP	27	-20.145	66.500	-42.647	1.00	32.70
	ATOM	209	CZ2	TRP	27	-19.551	66.020	-40.257	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-21.094	67.032	-38.706	1.00	26.93
	ATOM	211	CH2	TRP	27	-19.957	66.260	-38.975	1.00	28.08
	ATOM	212	C	TRP	27	-25.617	68.878	-42.192	1.00	38.50
	ATOM	213	O	TRP	27	-26.449	68.787	-43.088	1.00	39.07
	ATOM	214	N	LEU	28	-25.628	69.823	-41.259	1.00	40.19
20	ATOM	215	CA	LEU	28	-26.613	70.892	-41.207	1.00	39.45
	ATOM	216	CB	LEU	28	-27.624	70.626	-40.099	1.00	44.18
	ATOM	217	CG	LEU	28	-28.916	71.437	-40.170	1.00	46.83
	ATOM	218	CD1	LEU	28	-29.712	70.992	-41.400	1.00	45.15
	ATOM	219	CD2	LEU	28	-29.731	71.230	-38.890	1.00	49.15
25	ATOM	220	C	LEU	28	-25.793	72.127	-40.857	1.00	39.80
	ATOM	221	O	LEU	28	-24.898	72.051	-40.028	1.00	42.40
	ATOM	222	N	LEU	29	-26.077	73.257	-41.483	1.00	33.58
	ATOM	223	CA	LEU	29	-25.310	74.450	-41.181	1.00	29.21
	ATOM	224	CB	LEU	29	-25.689	75.595	-42.119	1.00	27.16
30	ATOM	225	CG	LEU	29	-24.970	76.913	-41.820	1.00	22.36
	ATOM	226	CD1	LEU	29	-23.521	76.816	-42.268	1.00	21.47
	ATOM	227	CD2	LEU	29	-25.662	78.057	-42.512	1.00	14.53
	ATOM	228	C	LEU	29	-25.498	74.916	-39.742	1.00	29.98
	ATOM	229	O	LEU	29	-26.616	75.105	-39.273	1.00	30.36
35	ATOM	230	N	SER	30	-24.383	75.095	-39.048	1.00	29.88
	ATOM	231	CA	SER	30	-24.406	75.581	-37.681	1.00	28.49
	ATOM	232	CB	SER	30	-23.199	75.062	-36.899	1.00	26.13
	ATOM	233	OG	SER	30	-23.173	75.611	-35.600	1.00	20.25
	ATOM	234	C	SER	30	-24.339	77.096	-37.820	1.00	31.61
40	ATOM	235	O	SER	30	-25.172	77.818	-37.280	1.00	35.14
	ATOM	236	N	PHE	31	-23.343	77.570	-38.561	1.00	32.12
	ATOM	237	CA	PHE	31	-23.183	78.993	-38.804	1.00	30.76
	ATOM	238	CB	PHE	31	-22.806	79.729	-37.512	1.00	27.55
	ATOM	239	CG	PHE	31	-21.328	79.789	-37.242	1.00	31.23
45	ATOM	240	CD1	PHE	31	-20.535	80.757	-37.848	1.00	32.00
	ATOM	241	CD2	PHE	31	-20.724	78.868	-36.390	1.00	27.89
	ATOM	242	CE1	PHE	31	-19.171	80.806	-37.612	1.00	29.70
	ATOM	243	CE2	PHE	31	-19.360	78.914	-36.150	1.00	26.15
	ATOM	244	CZ	PHE	31	-18.582	79.882	-36.761	1.00	28.30
50	ATOM	245	C	PHE	31	-22.123	79.219	-39.868	1.00	32.06
	ATOM	246	O	PHE	31	-21.200	78.423	-40.030	1.00	32.26
	ATOM	247	N	LYS	32	-22.277	80.306	-40.605	1.00	34.83
	ATOM	248	CA	LYS	32	-21.331	80.657	-41.644	1.00	37.13
	ATOM	249	CB	LYS	32	-21.969	80.474	-43.019	1.00	36.55
55	ATOM	250	CG	LYS	32	-21.071	80.882	-44.154	1.00	40.62
	ATOM	251	CD	LYS	32	-21.813	80.960	-45.464	1.00	44.56
	ATOM	252	CE	LYS	32	-20.935	81.666	-46.500	1.00	46.29
	ATOM	253	NZ	LYS	32	-21.536	81.680	-47.866	1.00	50.58
	ATOM	254	C	LYS	32	-20.950	82.118	-41.419	1.00	36.39
60	ATOM	255	O	LYS	32	-21.815	82.962	-41.187	1.00	38.69
	ATOM	256	N	ARG	33	-19.655	82.411	-41.467	1.00	37.90
	ATOM	257	CA	ARG	33	-19.184	83.773	-41.257	1.00	35.74
	ATOM	258	CB	ARG	33	-18.558	83.878	-39.871	1.00	36.79
	ATOM	259	CG	ARG	33	-17.948	85.219	-39.532	1.00	31.09
65	ATOM	260	CD	ARG	33	-17.615	85.268	-38.046	1.00	31.62
	ATOM	261	NE	ARG	33	-17.160	86.581	-37.596	1.00	38.16

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	ATOM	262	CZ	ARG	33	-15.887	86.901	-37.394	1.00	39.15
	ATOM	263	NH1	ARG	33	-14.935	86.002	-37.598	1.00	43.41
	ATOM	264	NH2	ARG	33	-15.563	88.118	-36.992	1.00	40.65
	ATOM	265	C	ARG	33	-18.170	84.148	-42.331	1.00	37.71
5	ATOM	266	O	ARG	33	-17.121	83.510	-42.460	1.00	41.16
	ATOM	267	N	GLY	34	-18.479	85.176	-43.113	1.00	36.59
	ATOM	268	CA	GLY	34	-17.560	85.585	-44.154	1.00	36.72
	ATOM	269	C	GLY	34	-17.855	84.929	-45.486	1.00	40.01
	ATOM	270	O	GLY	34	-18.917	84.341	-45.685	1.00	39.85
10	ATOM	271	N	SER	35	-16.891	85.009	-46.393	1.00	38.74
	ATOM	272	CA	SER	35	-17.050	84.459	-47.731	1.00	40.23
	ATOM	273	CB	SER	35	-16.917	85.592	-48.741	1.00	43.02
	ATOM	274	OG	SER	35	-15.695	86.300	-48.530	1.00	48.05
	ATOM	275	C	SER	35	-16.076	83.344	-48.119	1.00	40.54
15	ATOM	276	O	SER	35	-16.329	82.617	-49.078	1.00	43.74
	ATOM	277	N	ALA	36	-14.975	83.208	-47.384	1.00	35.56
	ATOM	278	CA	ALA	36	-13.958	82.208	-47.691	1.00	29.62
	ATOM	279	CB	ALA	36	-12.790	82.370	-46.741	1.00	25.47
	ATOM	280	C	ALA	36	-14.416	80.752	-47.701	1.00	31.23
20	ATOM	281	O	ALA	36	-13.808	79.924	-48.367	1.00	29.56
	ATOM	282	N	LEU	37	-15.480	80.434	-46.971	1.00	34.09
	ATOM	283	CA	LEU	37	-15.964	79.059	-46.900	1.00	32.32
	ATOM	284	CB	LEU	37	-15.636	78.467	-45.531	1.00	26.47
	ATOM	285	CG	LEU	37	-14.150	78.431	-45.185	1.00	25.72
25	ATOM	286	CD1	LEU	37	-13.959	78.284	-43.696	1.00	24.61
	ATOM	287	CD2	LEU	37	-13.481	77.302	-45.935	1.00	27.53
	ATOM	288	C	LEU	37	-17.455	78.938	-47.160	1.00	35.16
	ATOM	289	O	LEU	37	-18.242	79.780	-46.744	1.00	36.47
	ATOM	290	N	GLU	38	-17.833	77.871	-47.847	1.00	38.30
30	ATOM	291	CA	GLU	38	-19.227	77.615	-48.170	1.00	41.17
	ATOM	292	CB	GLU	38	-19.508	77.970	-49.626	1.00	46.15
	ATOM	293	CG	GLU	38	-19.915	79.402	-49.882	1.00	50.01
	ATOM	294	CD	GLU	38	-19.900	79.731	-51.369	1.00	51.18
	ATOM	295	OE1	GLU	38	-20.320	78.858	-52.171	1.00	48.40
35	ATOM	296	OE2	GLU	38	-19.470	80.858	-51.729	1.00	50.23
	ATOM	297	C	GLU	38	-19.573	76.151	-47.977	1.00	42.69
	ATOM	298	O	GLU	38	-18.707	75.287	-48.007	1.00	42.55
	ATOM	299	N	GLU	39	-20.850	75.870	-47.781	1.00	44.75
	ATOM	300	CA	GLU	39	-21.277	74.489	-47.628	1.00	46.50
40	ATOM	301	CB	GLU	39	-22.441	74.391	-46.647	1.00	47.52
	ATOM	302	CG	GLU	39	-23.422	75.536	-46.781	1.00	58.50
	ATOM	303	CD	GLU	39	-24.645	75.359	-45.902	1.00	62.35
	ATOM	304	OE1	GLU	39	-25.436	76.330	-45.774	1.00	64.58
	ATOM	305	OE2	GLU	39	-24.818	74.248	-45.349	1.00	63.04
45	ATOM	306	C	GLU	39	-21.721	74.087	-49.022	1.00	44.56
	ATOM	307	O	GLU	39	-22.371	74.860	-49.714	1.00	48.11
	ATOM	308	N	LYS	40	-21.355	72.888	-49.447	1.00	40.71
	ATOM	309	CA	LYS	40	-21.724	72.426	-50.770	1.00	38.08
	ATOM	310	CB	LYS	40	-20.682	72.874	-51.794	1.00	41.28
50	ATOM	311	CG	LYS	40	-20.886	72.286	-53.185	1.00	41.52
	ATOM	312	CD	LYS	40	-19.652	72.497	-54.051	1.00	42.73
	ATOM	313	CE	LYS	40	-19.761	71.785	-55.388	1.00	41.97
	ATOM	314	NZ	LYS	40	-18.487	71.907	-56.162	1.00	45.94
	ATOM	315	C	LYS	40	-21.845	70.917	-50.808	1.00	39.66
55	ATOM	316	O	LYS	40	-20.861	70.200	-50.649	1.00	36.18
	ATOM	317	N	GLU	41	-23.066	70.445	-51.019	1.00	40.16
	ATOM	318	CA	GLU	41	-23.341	69.019	-51.106	1.00	42.80
	ATOM	319	CB	GLU	41	-22.864	68.508	-52.461	1.00	45.31
	ATOM	320	CG	GLU	41	-23.476	69.297	-53.612	1.00	56.07
60	ATOM	321	CD	GLU	41	-22.728	69.124	-54.931	1.00	59.42
	ATOM	322	OE1	GLU	41	-21.496	69.366	-54.969	1.00	63.71
	ATOM	323	OE2	GLU	41	-23.376	68.751	-55.934	1.00	63.71
	ATOM	324	C	GLU	41	-22.701	68.233	-49.974	1.00	38.93
	ATOM	325	O	GLU	41	-21.967	67.276	-50.205	1.00	39.03
65	ATOM	326	N	ASN	42	-22.976	68.665	-48.751	1.00	34.69
	ATOM	327	CA	ASN	42	-22.474	68.011	-47.555	1.00	32.96

	ATOM	328	CB	ASN	42	-22.866	66.538	-47.562	1.00	32.65
	ATOM	329	CG	ASN	42	-23.157	66.024	-46.187	1.00	35.50
	ATOM	330	OD1	ASN	42	-22.732	64.937	-45.819	1.00	40.26
	ATOM	331	ND2	ASN	42	-23.896	66.804	-45.413	1.00	29.28
5	ATOM	332	C	ASN	42	-20.973	68.118	-47.333	1.00	31.44
	ATOM	333	O	ASN	42	-20.401	67.359	-46.558	1.00	22.78
	ATOM	334	N	LYS	43	-20.342	69.066	-48.008	1.00	33.26
	ATOM	335	CA	LYS	43	-18.908	69.264	-47.876	1.00	33.68
	ATOM	336	CB	LYS	43	-18.186	68.744	-49.119	1.00	33.55
10	ATOM	337	CG	LYS	43	-18.254	67.247	-49.281	1.00	40.49
	ATOM	338	CD	LYS	43	-17.606	66.814	-50.572	1.00	45.39
	ATOM	339	CE	LYS	43	-18.381	67.338	-51.764	1.00	52.57
	ATOM	340	NZ	LYS	43	-17.814	66.830	-53.040	1.00	62.06
	ATOM	341	C	LYS	43	-18.614	70.737	-47.706	1.00	32.05
15	ATOM	342	O	LYS	43	-19.476	71.573	-47.941	1.00	37.32
	ATOM	343	N	ILE	44	-17.402	71.060	-47.281	1.00	31.31
	ATOM	344	CA	ILE	44	-17.045	72.455	-47.122	1.00	29.52
	ATOM	345	CB	ILE	44	-16.247	72.687	-45.839	1.00	26.38
	ATOM	346	CG2	ILE	44	-15.855	74.146	-45.737	1.00	26.09
20	ATOM	347	CG1	ILE	44	-17.103	72.303	-44.631	1.00	28.61
	ATOM	348	CD1	ILE	44	-16.435	72.510	-43.307	1.00	21.51
	ATOM	349	C	ILE	44	-16.230	72.888	-48.328	1.00	31.60
	ATOM	350	O	ILE	44	-15.193	72.302	-48.635	1.00	34.50
	ATOM	351	N	LEU	45	-16.721	73.904	-49.023	1.00	31.44
25	ATOM	352	CA	LEU	45	-16.053	74.427	-50.205	1.00	31.40
	ATOM	353	CB	LEU	45	-17.090	74.759	-51.275	1.00	33.60
	ATOM	354	CG	LEU	45	-16.558	75.428	-52.540	1.00	34.36
	ATOM	355	CD1	LEU	45	-15.669	74.470	-53.305	1.00	29.00
	ATOM	356	CD2	LEU	45	-17.733	75.869	-53.392	1.00	32.42
30	ATOM	357	C	LEU	45	-15.222	75.670	-49.898	1.00	27.80
	ATOM	358	O	LEU	45	-15.702	76.620	-49.289	1.00	27.94
	ATOM	359	N	VAL	46	-13.967	75.649	-50.330	1.00	28.00
	ATOM	360	CA	VAL	46	-13.058	76.767	-50.124	1.00	31.53
	ATOM	361	CB	VAL	46	-11.606	76.263	-50.070	1.00	27.89
35	ATOM	362	CG1	VAL	46	-10.660	77.409	-49.806	1.00	26.10
	ATOM	363	CG2	VAL	46	-11.477	75.214	-49.001	1.00	23.39
	ATOM	364	C	VAL	46	-13.211	77.771	-51.278	1.00	33.94
	ATOM	365	O	VAL	46	-13.072	77.405	-52.447	1.00	39.68
	ATOM	366	N	LYS	47	-13.492	79.030	-50.950	1.00	33.59
40	ATOM	367	CA	LYS	47	-13.666	80.058	-51.974	1.00	37.05
	ATOM	368	CB	LYS	47	-14.963	80.836	-51.727	1.00	34.49
	ATOM	369	CG	LYS	47	-16.171	79.950	-51.482	1.00	37.53
	ATOM	370	CD	LYS	47	-16.407	78.988	-52.639	1.00	41.69
	ATOM	371	CE	LYS	47	-17.413	79.548	-53.627	1.00	42.95
45	ATOM	372	NZ	LYS	47	-17.115	80.958	-53.966	1.00	44.50
	ATOM	373	C	LYS	47	-12.491	81.032	-52.065	1.00	37.34
	ATOM	374	O	LYS	47	-12.398	81.809	-53.010	1.00	43.04
	ATOM	375	N	GLU	48	-11.605	80.987	-51.081	1.00	36.99
	ATOM	376	CA	GLU	48	-10.423	81.845	-51.052	1.00	37.31
50	ATOM	377	CB	GLU	48	-10.590	82.970	-50.044	1.00	36.53
	ATOM	378	CG	GLU	48	-11.609	84.010	-50.403	1.00	46.74
	ATOM	379	CD	GLU	48	-11.734	85.061	-49.315	1.00	52.39
	ATOM	380	OE1	GLU	48	-10.678	85.460	-48.751	1.00	51.33
	ATOM	381	OE2	GLU	48	-12.882	85.488	-49.027	1.00	54.55
55	ATOM	382	C	GLU	48	-9.268	80.988	-50.598	1.00	36.94
	ATOM	383	O	GLU	48	-9.372	80.326	-49.574	1.00	40.21
	ATOM	384	N	THR	49	-8.158	80.994	-51.326	1.00	36.34
	ATOM	385	CA	THR	49	-7.052	80.153	-50.897	1.00	35.48
	ATOM	386	CB	THR	49	-6.058	79.862	-52.056	1.00	31.16
60	ATOM	387	OG1	THR	49	-4.955	80.762	-51.981	1.00	32.48
	ATOM	388	CG2	THR	49	-6.736	80.021	-53.396	1.00	30.36
	ATOM	389	C	THR	49	-6.320	80.790	-49.713	1.00	34.59
	ATOM	390	O	THR	49	-6.296	82.010	-49.562	1.00	35.13
	ATOM	391	N	GLY	50	-5.745	79.942	-48.863	1.00	32.49
65	ATOM	392	CA	GLY	50	-5.025	80.426	-47.705	1.00	28.17
	ATOM	393	C	GLY	50	-4.814	79.317	-46.700	1.00	29.99

	ATOM	394	O	GLY	50	-4.915	78.148	-47.038	1.00	29.76
	ATOM	395	N	TYR	51	-4.508	79.687	-45.461	1.00	31.68
	ATOM	396	CA	TYR	51	-4.298	78.718	-44.391	1.00	32.88
	ATOM	397	CB	TYR	51	-3.076	79.099	-43.548	1.00	35.97
5	ATOM	398	CG	TYR	51	-1.780	78.913	-44.280	1.00	43.03
	ATOM	399	CD1	TYR	51	-1.319	79.877	-45.182	1.00	44.15
	ATOM	400	CE1	TYR	51	-0.163	79.667	-45.934	1.00	47.81
	ATOM	401	CD2	TYR	51	-1.052	77.735	-44.139	1.00	47.08
	ATOM	402	CE2	TYR	51	0.103	77.507	-44.886	1.00	51.79
10	ATOM	403	CZ	TYR	51	0.543	78.475	-45.786	1.00	51.15
	ATOM	404	OH	TYR	51	1.659	78.226	-46.563	1.00	52.20
	ATOM	405	C	TYR	51	-5.529	78.621	-43.500	1.00	31.97
	ATOM	406	O	TYR	51	-6.022	79.623	-42.988	1.00	30.01
	ATOM	407	N	PHE	52	-6.025	77.402	-43.317	1.00	31.50
15	ATOM	408	CA	PHE	52	-7.207	77.195	-42.497	1.00	30.41
	ATOM	409	CB	PHE	52	-8.357	76.645	-43.343	1.00	31.30
	ATOM	410	CG	PHE	52	-8.763	77.528	-44.481	1.00	29.70
	ATOM	411	CD1	PHE	52	-7.967	77.642	-45.609	1.00	26.20
	ATOM	412	CD2	PHE	52	-9.952	78.240	-44.427	1.00	28.38
20	ATOM	413	CE1	PHE	52	-8.350	78.449	-46.660	1.00	27.17
	ATOM	414	CE2	PHE	52	-10.342	79.051	-45.476	1.00	27.76
	ATOM	415	CZ	PHE	52	-9.542	79.156	-46.594	1.00	27.27
	ATOM	416	C	PHE	52	-6.985	76.240	-41.343	1.00	27.65
	ATOM	417	O	PHE	52	-6.205	75.301	-41.438	1.00	24.85
25	ATOM	418	N	PHE	53	-7.682	76.504	-40.248	1.00	27.18
	ATOM	419	CA	PHE	53	-7.646	75.644	-39.075	1.00	27.05
	ATOM	420	CB	PHE	53	-7.783	76.461	-37.797	1.00	28.76
	ATOM	421	CG	PHE	53	-7.947	75.627	-36.567	1.00	28.00
	ATOM	422	CD1	PHE	53	-6.889	74.884	-36.066	1.00	26.49
30	ATOM	423	CD2	PHE	53	-9.170	75.567	-35.915	1.00	28.04
	ATOM	424	CE1	PHE	53	-7.047	74.096	-34.932	1.00	26.54
	ATOM	425	CE2	PHE	53	-9.332	74.778	-34.780	1.00	25.78
	ATOM	426	CZ	PHE	53	-8.268	74.044	-34.291	1.00	24.39
	ATOM	427	C	PHE	53	-8.892	74.785	-39.269	1.00	27.56
35	ATOM	428	O	PHE	53	-9.997	75.312	-39.360	1.00	25.39
	ATOM	429	N	ILE	54	-8.710	73.471	-39.350	1.00	28.87
	ATOM	430	CA	ILE	54	-9.823	72.554	-39.575	1.00	28.95
	ATOM	431	CB	ILE	54	-9.565	71.726	-40.850	1.00	29.60
	ATOM	432	CG2	ILE	54	-10.827	70.977	-41.247	1.00	27.42
40	ATOM	433	CG1	ILE	54	-9.131	72.657	-41.984	1.00	26.41
	ATOM	434	CD1	ILE	54	-8.538	71.962	-43.170	1.00	28.70
	ATOM	435	C	ILE	54	-10.028	71.621	-38.382	1.00	28.57
	ATOM	436	O	ILE	54	-9.073	71.095	-37.821	1.00	29.17
	ATOM	437	N	TYR	55	-11.279	71.407	-37.995	1.00	26.97
45	ATOM	438	CA	TYR	55	-11.558	70.554	-36.846	1.00	24.22
	ATOM	439	CB	TYR	55	-11.724	71.419	-35.596	1.00	21.82
	ATOM	440	CG	TYR	55	-12.824	72.446	-35.694	1.00	27.68
	ATOM	441	CD1	TYR	55	-14.122	72.148	-35.289	1.00	27.01
	ATOM	442	CE1	TYR	55	-15.135	73.084	-35.399	1.00	27.44
50	ATOM	443	CD2	TYR	55	-12.573	73.712	-36.210	1.00	27.16
	ATOM	444	CE2	TYR	55	-13.583	74.654	-36.326	1.00	25.39
	ATOM	445	CZ	TYR	55	-14.858	74.334	-35.921	1.00	29.02
	ATOM	446	OH	TYR	55	-15.858	75.263	-36.047	1.00	31.82
	ATOM	447	C	TYR	55	-12.776	69.677	-37.036	1.00	25.16
55	ATOM	448	O	TYR	55	-13.583	69.916	-37.906	1.00	28.47
	ATOM	449	N	GLY	56	-12.900	68.644	-36.219	1.00	27.44
	ATOM	450	CA	GLY	56	-14.036	67.757	-36.337	1.00	22.07
	ATOM	451	C	GLY	56	-14.142	66.803	-35.168	1.00	25.83
	ATOM	452	O	GLY	56	-13.133	66.386	-34.608	1.00	26.00
60	ATOM	453	N	GLN	57	-15.370	66.479	-34.785	1.00	22.56
	ATOM	454	CA	GLN	57	-15.625	65.552	-33.694	1.00	20.91
	ATOM	455	CB	GLN	57	-15.894	66.283	-32.378	1.00	17.87
	ATOM	456	CG	GLN	57	-16.294	65.336	-31.253	1.00	16.86
	ATOM	457	CD	GLN	57	-16.485	66.022	-29.917	1.00	22.06
65	ATOM	458	OE1	GLN	57	-15.545	66.563	-29.346	1.00	27.26
	ATOM	459	NE2	GLN	57	-17.711	65.997	-29.408	1.00	20.20

	ATOM	460	C	GLN	57	-16.828	64.701	-34.042	1.00	24.23
	ATOM	461	O	GLN	57	-17.768	65.179	-34.675	1.00	25.16
	ATOM	462	N	VAL	58	-16.784	63.440	-33.623	1.00	24.62
	ATOM	463	CA	VAL	58	-17.858	62.483	-33.863	1.00	26.68
5	ATOM	464	CB	VAL	58	-17.517	61.533	-35.036	1.00	25.55
	ATOM	465	CG1	VAL	58	-18.541	60.428	-35.127	1.00	15.69
	ATOM	466	CG2	VAL	58	-17.467	62.300	-36.340	1.00	25.77
	ATOM	467	C	VAL	58	-18.009	61.639	-32.614	1.00	31.06
	ATOM	468	O	VAL	58	-17.011	61.284	-32.001	1.00	27.74
10	ATOM	469	N	LEU	59	-19.247	61.323	-32.232	1.00	33.82
	ATOM	470	CA	LEU	59	-19.501	60.471	-31.065	1.00	29.85
	ATOM	471	CB	LEU	59	-20.707	60.961	-30.269	1.00	29.62
	ATOM	472	CG	LEU	59	-20.787	60.557	-28.790	1.00	28.11
	ATOM	473	CD1	LEU	59	-22.188	60.822	-28.278	1.00	26.67
15	ATOM	474	CD2	LEU	59	-20.441	59.110	-28.595	1.00	22.45
	ATOM	475	C	LEU	59	-19.789	59.060	-31.572	1.00	30.98
	ATOM	476	O	LEU	59	-20.789	58.826	-32.247	1.00	32.52
	ATOM	477	N	TYR	60	-18.905	58.122	-31.248	1.00	33.01
	ATOM	478	CA	TYR	60	-19.079	56.743	-31.680	1.00	34.51
20	ATOM	479	CB	TYR	60	-17.721	56.111	-31.971	1.00	40.10
	ATOM	480	CG	TYR	60	-16.992	56.804	-33.080	1.00	45.37
	ATOM	481	CD1	TYR	60	-15.901	57.634	-32.817	1.00	47.13
	ATOM	482	CE1	TYR	60	-15.283	58.341	-33.838	1.00	50.28
	ATOM	483	CD2	TYR	60	-17.440	56.695	-34.390	1.00	44.35
25	ATOM	484	CE2	TYR	60	-16.836	57.393	-35.417	1.00	48.81
	ATOM	485	CZ	TYR	60	-15.764	58.216	-35.137	1.00	51.91
	ATOM	486	OH	TYR	60	-15.200	58.939	-36.159	1.00	53.77
	ATOM	487	C	TYR	60	-19.830	55.889	-30.673	1.00	35.49
	ATOM	488	O	TYR	60	-19.430	55.762	-29.521	1.00	36.13
30	ATOM	489	N	THR	61	-20.930	55.306	-31.127	1.00	36.19
	ATOM	490	CA	THR	61	-21.744	54.445	-30.290	1.00	35.36
	ATOM	491	CB	THR	61	-23.156	55.012	-30.135	1.00	34.22
	ATOM	492	OG1	THR	61	-23.709	55.272	-31.429	1.00	34.22
	ATOM	493	CG2	THR	61	-23.127	56.297	-29.330	1.00	25.45
35	ATOM	494	C	THR	61	-21.798	53.088	-30.978	1.00	37.82
	ATOM	495	O	THR	61	-22.789	52.369	-30.907	1.00	43.70
	ATOM	496	N	ASP	62	-20.703	52.762	-31.652	1.00	37.64
	ATOM	497	CA	ASP	62	-20.543	51.522	-32.390	1.00	38.08
	ATOM	498	CB	ASP	62	-20.011	51.867	-33.777	1.00	38.42
40	ATOM	499	CG	ASP	62	-19.974	50.687	-34.704	1.00	41.25
	ATOM	500	OD1	ASP	62	-20.290	50.874	-35.903	1.00	38.28
	ATOM	501	OD2	ASP	62	-19.619	49.586	-34.235	1.00	41.37
	ATOM	502	C	ASP	62	-19.544	50.664	-31.619	1.00	40.90
	ATOM	503	O	ASP	62	-18.579	51.191	-31.084	1.00	46.73
45	ATOM	504	N	LYS	63	-19.756	49.352	-31.549	1.00	42.62
	ATOM	505	CA	LYS	63	-18.825	48.510	-30.797	1.00	40.92
	ATOM	506	CB	LYS	63	-19.593	47.488	-29.957	1.00	42.79
	ATOM	507	CG	LYS	63	-20.491	46.560	-30.744	1.00	45.91
	ATOM	508	CD	LYS	63	-21.365	45.737	-29.798	1.00	46.22
50	ATOM	509	CE	LYS	63	-22.288	46.647	-28.964	1.00	49.97
	ATOM	510	NZ	LYS	63	-23.013	45.933	-27.857	1.00	49.98
	ATOM	511	C	LYS	63	-17.767	47.794	-31.631	1.00	42.59
	ATOM	512	O	LYS	63	-17.169	46.827	-31.176	1.00	42.46
	ATOM	513	N	THR	64	-17.512	48.302	-32.830	1.00	44.06
55	ATOM	514	CA	THR	64	-16.535	47.714	-33.735	1.00	44.94
	ATOM	515	CB	THR	64	-16.675	48.333	-35.148	1.00	43.79
	ATOM	516	OG1	THR	64	-18.025	48.180	-35.598	1.00	46.20
	ATOM	517	CG2	THR	64	-15.755	47.641	-36.147	1.00	42.80
	ATOM	518	C	THR	64	-15.062	47.799	-33.299	1.00	45.82
60	ATOM	519	O	THR	64	-14.201	48.108	-34.102	1.00	53.10
	ATOM	520	N	TYR	65	-14.771	47.516	-32.039	1.00	45.11
	ATOM	521	CA	TYR	65	-13.398	47.522	-31.512	1.00	45.72
	ATOM	522	CB	TYR	65	-12.648	46.260	-31.979	1.00	40.46
	ATOM	523	CG	TYR	65	-11.817	46.412	-33.233	1.00	41.90
65	ATOM	524	CD1	TYR	65	-10.493	46.853	-33.173	1.00	42.47
	ATOM	525	CE1	TYR	65	-9.719	47.005	-34.337	1.00	41.83

	ATOM	526	CD2	TYR	65	-12.354	46.123	-34.492	1.00	41.65
	ATOM	527	CE2	TYR	65	-11.593	46.273	-35.665	1.00	40.28
	ATOM	528	CZ	TYR	65	-10.280	46.714	-35.575	1.00	41.90
	ATOM	529	OH	TYR	65	-9.532	46.863	-36.717	1.00	43.84
5	ATOM	530	C	TYR	65	-12.531	48.763	-31.776	1.00	45.31
	ATOM	531	O	TYR	65	-11.659	49.104	-30.965	1.00	49.74
	ATOM	532	N	ALA	66	-12.753	49.432	-32.899	1.00	42.32
	ATOM	533	CA	ALA	66	-11.986	50.622	-33.244	1.00	39.57
	ATOM	534	CB	ALA	66	-10.651	50.230	-33.839	1.00	35.36
10	ATOM	535	C	ALA	66	-12.769	51.468	-34.232	1.00	39.85
	ATOM	536	O	ALA	66	-13.135	51.004	-35.313	1.00	37.60
	ATOM	537	N	MET	67	-13.042	52.708	-33.847	1.00	41.88
	ATOM	538	CA	MET	67	-13.780	53.626	-34.702	1.00	41.48
	ATOM	539	CB	MET	67	-15.135	53.984	-34.080	1.00	39.98
15	ATOM	540	CG	MET	67	-16.139	52.839	-34.029	1.00	37.31
	ATOM	541	SD	MET	67	-16.527	52.168	-35.651	1.00	35.46
	ATOM	542	CE	MET	67	-17.731	53.352	-36.258	1.00	34.06
	ATOM	543	C	MET	67	-12.958	54.886	-34.909	1.00	39.50
	ATOM	544	O	MET	67	-11.976	55.124	-34.202	1.00	38.44
20	ATOM	545	N	GLY	68	-13.364	55.689	-35.884	1.00	39.84
	ATOM	546	CA	GLY	68	-12.653	56.920	-36.165	1.00	37.61
	ATOM	547	C	GLY	68	-13.120	57.566	-37.447	1.00	33.76
	ATOM	548	O	GLY	68	-13.909	56.987	-38.185	1.00	35.58
	ATOM	549	N	HIS	69	-12.644	58.776	-37.708	1.00	35.07
25	ATOM	550	CA	HIS	69	-13.019	59.485	-38.920	1.00	33.70
	ATOM	551	CB	HIS	69	-14.098	60.544	-38.627	1.00	37.02
	ATOM	552	CG	HIS	69	-13.747	61.502	-37.531	1.00	34.65
	ATOM	553	CD2	HIS	69	-13.370	62.800	-37.572	1.00	35.63
	ATOM	554	ND1	HIS	69	-13.781	61.158	-36.197	1.00	37.68
30	ATOM	555	CE1	HIS	69	-13.442	62.202	-35.463	1.00	32.64
	ATOM	556	NE2	HIS	69	-13.188	63.212	-36.274	1.00	35.90
	ATOM	557	C	HIS	69	-11.813	60.137	-39.582	1.00	33.98
	ATOM	558	O	HIS	69	-10.738	60.195	-39.007	1.00	30.94
	ATOM	559	N	LEU	70	-12.003	60.622	-40.801	1.00	35.35
35	ATOM	560	CA	LEU	70	-10.935	61.271	-41.546	1.00	30.76
	ATOM	561	CB	LEU	70	-10.576	60.450	-42.780	1.00	29.59
	ATOM	562	CG	LEU	70	-10.530	58.929	-42.673	1.00	30.71
	ATOM	563	CD1	LEU	70	-10.413	58.339	-44.057	1.00	25.60
	ATOM	564	CD2	LEU	70	-9.373	58.507	-41.798	1.00	27.39
40	ATOM	565	C	LEU	70	-11.390	62.644	-42.031	1.00	33.19
	ATOM	566	O	LEU	70	-12.526	62.805	-42.478	1.00	37.07
	ATOM	567	N	ILE	71	-10.520	63.641	-41.922	1.00	30.93
	ATOM	568	CA	ILE	71	-10.852	64.958	-42.443	1.00	27.94
	ATOM	569	CB	ILE	71	-10.364	66.086	-41.530	1.00	28.22
45	ATOM	570	CG2	ILE	71	-10.420	67.411	-42.263	1.00	24.54
	ATOM	571	CG1	ILE	71	-11.246	66.154	-40.290	1.00	26.34
	ATOM	572	CD1	ILE	71	-10.735	67.097	-39.237	1.00	37.50
	ATOM	573	C	ILE	71	-10.061	64.950	-43.741	1.00	30.32
	ATOM	574	O	ILE	71	-8.837	64.885	-43.716	1.00	23.07
50	ATOM	575	N	GLN	72	-10.757	64.984	-44.874	1.00	29.74
	ATOM	576	CA	GLN	72	-10.070	64.927	-46.155	1.00	27.85
	ATOM	577	CB	GLN	72	-10.582	63.731	-46.943	1.00	26.70
	ATOM	578	CG	GLN	72	-10.568	62.463	-46.138	1.00	28.28
	ATOM	579	CD	GLN	72	-10.883	61.253	-46.970	1.00	33.11
55	ATOM	580	OE1	GLN	72	-11.903	61.210	-47.651	1.00	41.22
	ATOM	581	NE2	GLN	72	-10.011	60.255	-46.919	1.00	30.18
	ATOM	582	C	GLN	72	-10.166	66.172	-47.011	1.00	29.85
	ATOM	583	O	GLN	72	-11.081	66.984	-46.860	1.00	28.20
	ATOM	584	N	ARG	73	-9.212	66.297	-47.927	1.00	30.90
60	ATOM	585	CA	ARG	73	-9.153	67.427	-48.839	1.00	30.79
	ATOM	586	CB	ARG	73	-7.889	68.230	-48.567	1.00	29.18
	ATOM	587	CG	ARG	73	-7.678	69.373	-49.513	1.00	28.66
	ATOM	588	CD	ARG	73	-6.226	69.743	-49.561	1.00	29.25
	ATOM	589	NE	ARG	73	-5.991	70.885	-50.431	1.00	35.78
65	ATOM	590	CZ	ARG	73	-4.808	71.207	-50.932	1.00	35.27
	ATOM	591	NH1	ARG	73	-3.744	70.467	-50.654	1.00	34.95

	ATOM	592	NH2	ARG	73	-4.691	72.271	-51.705	1.00	33.33
	ATOM	593	C	ARG	73	-9.154	66.980	-50.299	1.00	32.42
	ATOM	594	O	ARG	73	-8.360	66.131	-50.689	1.00	34.19
	ATOM	595	N	LYS	74	-10.062	67.540	-51.093	1.00	35.90
5	ATOM	596	CA	LYS	74	-10.129	67.235	-52.522	1.00	38.57
	ATOM	597	CB	LYS	74	-11.578	67.100	-52.998	1.00	43.52
	ATOM	598	CG	LYS	74	-12.282	65.831	-52.549	1.00	52.89
	ATOM	599	CD	LYS	74	-13.753	65.779	-53.022	1.00	58.14
	ATOM	600	CE	LYS	74	-14.438	64.498	-52.542	1.00	56.75
10	ATOM	601	NZ	LYS	74	-15.874	64.439	-52.946	1.00	63.98
	ATOM	602	C	LYS	74	-9.485	68.402	-53.260	1.00	40.11
	ATOM	603	O	LYS	74	-10.078	69.484	-53.357	1.00	37.55
	ATOM	604	N	LYS	75	-8.279	68.186	-53.775	1.00	34.48
	ATOM	605	CA	LYS	75	-7.555	69.229	-54.493	1.00	34.24
15	ATOM	606	CB	LYS	75	-6.127	68.764	-54.786	1.00	36.16
	ATOM	607	CG	LYS	75	-5.281	68.355	-53.582	1.00	35.99
	ATOM	608	CD	LYS	75	-3.894	67.905	-54.060	1.00	40.91
	ATOM	609	CE	LYS	75	-2.989	67.507	-52.902	1.00	46.16
	ATOM	610	NZ	LYS	75	-1.632	67.003	-53.313	1.00	46.03
20	ATOM	611	C	LYS	75	-8.239	69.579	-55.818	1.00	30.92
	ATOM	612	O	LYS	75	-8.713	68.700	-56.514	1.00	27.47
	ATOM	613	N	VAL	76	-8.279	70.864	-56.161	1.00	32.82
	ATOM	614	CA	VAL	76	-8.874	71.313	-57.423	1.00	31.51
	ATOM	615	CB	VAL	76	-9.156	72.826	-57.452	1.00	29.19
25	ATOM	616	CG1	VAL	76	-10.403	73.095	-58.244	1.00	27.09
	ATOM	617	CG2	VAL	76	-9.261	73.371	-56.072	1.00	34.94
	ATOM	618	C	VAL	76	-7.853	71.078	-58.517	1.00	33.40
	ATOM	619	O	VAL	76	-8.190	70.665	-59.621	1.00	32.92
	ATOM	620	N	HIS	77	-6.598	71.373	-58.192	1.00	34.88
30	ATOM	621	CA	HIS	77	-5.491	71.216	-59.117	1.00	34.59
	ATOM	622	CB	HIS	77	-4.623	72.479	-59.117	1.00	35.92
	ATOM	623	CG	HIS	77	-5.370	73.727	-59.474	1.00	34.17
	ATOM	624	CD2	HIS	77	-5.134	75.024	-59.171	1.00	31.44
	ATOM	625	ND1	HIS	77	-6.499	73.721	-60.268	1.00	35.05
35	ATOM	626	CE1	HIS	77	-6.929	74.957	-60.436	1.00	29.84
	ATOM	627	NE2	HIS	77	-6.116	75.766	-59.783	1.00	38.35
	ATOM	628	C	HIS	77	-4.669	70.005	-58.711	1.00	35.63
	ATOM	629	O	HIS	77	-4.430	69.785	-57.523	1.00	40.24
	ATOM	630	N	VAL	78	-4.216	69.231	-59.696	1.00	37.37
40	ATOM	631	CA	VAL	78	-3.471	68.023	-59.392	1.00	41.32
	ATOM	632	CB	VAL	78	-4.254	66.792	-59.887	1.00	43.10
	ATOM	633	CG1	VAL	78	-3.560	65.525	-59.449	1.00	45.01
	ATOM	634	CG2	VAL	78	-5.657	66.808	-59.306	1.00	44.98
	ATOM	635	C	VAL	78	-1.999	67.895	-59.804	1.00	43.02
45	ATOM	636	O	VAL	78	-1.135	67.839	-58.931	1.00	49.79
	ATOM	637	N	PHE	79	-1.680	67.831	-61.091	1.00	41.33
	ATOM	638	CA	PHE	79	-0.261	67.687	-61.496	1.00	45.00
	ATOM	639	CB	PHE	79	0.694	68.570	-60.668	1.00	40.51
	ATOM	640	CG	PHE	79	0.347	70.020	-60.662	1.00	39.52
50	ATOM	641	CD1	PHE	79	-0.231	70.599	-59.538	1.00	36.80
	ATOM	642	CD2	PHE	79	0.603	70.811	-61.774	1.00	37.89
	ATOM	643	CE1	PHE	79	-0.550	71.947	-59.521	1.00	37.52
	ATOM	644	CE2	PHE	79	0.288	72.160	-61.764	1.00	35.59
	ATOM	645	CZ	PHE	79	-0.291	72.731	-60.636	1.00	37.58
55	ATOM	646	C	PHE	79	0.295	66.268	-61.372	1.00	43.50
	ATOM	647	O	PHE	79	0.269	65.666	-60.297	1.00	37.85
	ATOM	648	N	GLY	80	0.832	65.762	-62.476	1.00	46.65
	ATOM	649	CA	GLY	80	1.433	64.443	-62.497	1.00	48.38
	ATOM	650	C	GLY	80	0.665	63.313	-61.848	1.00	47.89
60	ATOM	651	O	GLY	80	-0.530	63.131	-62.098	1.00	51.15
	ATOM	652	N	ASP	81	1.360	62.549	-61.008	1.00	45.07
	ATOM	653	CA	ASP	81	0.759	61.409	-60.341	1.00	43.99
	ATOM	654	CB	ASP	81	1.749	60.237	-60.309	1.00	45.66
	ATOM	655	CG	ASP	81	2.933	60.486	-59.389	1.00	47.49
65	ATOM	656	OD1	ASP	81	3.153	61.645	-58.971	1.00	51.90
	ATOM	657	OD2	ASP	81	3.658	59.518	-59.087	1.00	48.08

	ATOM	658	C	ASP	81	0.254	61.701	-58.939	1.00	42.29
	ATOM	659	O	ASP	81	0.126	60.789	-58.123	1.00	39.94
	ATOM	660	N	GLU	82	-0.032	62.968	-58.656	1.00	39.91
	ATOM	661	CA	GLU	82	-0.558	63.345	-57.346	1.00	36.58
5	ATOM	662	CB	GLU	82	-0.756	64.857	-57.227	1.00	34.91
	ATOM	663	CG	GLU	82	0.454	65.692	-56.955	1.00	37.82
	ATOM	664	CD	GLU	82	0.079	67.003	-56.284	1.00	40.75
	ATOM	665	OE1	GLU	82	-0.907	67.636	-56.710	1.00	38.15
	ATOM	666	OE2	GLU	82	0.768	67.406	-55.323	1.00	47.99
10	ATOM	667	C	GLU	82	-1.933	62.720	-57.160	1.00	36.58
	ATOM	668	O	GLU	82	-2.672	62.531	-58.123	1.00	38.40
	ATOM	669	N	LEU	83	-2.273	62.401	-55.919	1.00	33.83
	ATOM	670	CA	LEU	83	-3.592	61.874	-55.606	1.00	32.70
	ATOM	671	CB	LEU	83	-3.546	61.008	-54.347	1.00	30.78
15	ATOM	672	CG	LEU	83	-3.344	59.504	-54.519	1.00	29.26
	ATOM	673	CD1	LEU	83	-2.284	59.218	-55.553	1.00	29.60
	ATOM	674	CD2	LEU	83	-2.968	58.906	-53.188	1.00	28.65
	ATOM	675	C	LEU	83	-4.367	63.144	-55.328	1.00	33.88
	ATOM	676	O	LEU	83	-3.883	64.008	-54.606	1.00	38.46
20	ATOM	677	N	SER	84	-5.554	63.277	-55.902	1.00	34.94
	ATOM	678	CA	SER	84	-6.332	64.488	-55.689	1.00	39.77
	ATOM	679	CB	SER	84	-7.315	64.699	-56.839	1.00	38.92
	ATOM	680	OG	SER	84	-8.182	63.597	-56.954	1.00	43.11
	ATOM	681	C	SER	84	-7.076	64.501	-54.357	1.00	39.89
25	ATOM	682	O	SER	84	-7.606	65.530	-53.937	1.00	42.52
	ATOM	683	N	LEU	85	-7.108	63.356	-53.692	1.00	40.92
	ATOM	684	CA	LEU	85	-7.772	63.246	-52.407	1.00	34.66
	ATOM	685	CB	LEU	85	-8.755	62.075	-52.427	1.00	35.12
	ATOM	686	CG	LEU	85	-9.792	61.878	-51.313	1.00	36.92
30	ATOM	687	CD1	LEU	85	-9.108	61.508	-50.019	1.00	38.24
	ATOM	688	CD2	LEU	85	-10.619	63.145	-51.149	1.00	36.08
	ATOM	689	C	LEU	85	-6.693	63.020	-51.365	1.00	35.30
	ATOM	690	O	LEU	85	-5.995	62.010	-51.380	1.00	32.85
	ATOM	691	N	VAL	86	-6.542	63.984	-50.470	1.00	33.26
35	ATOM	692	CA	VAL	86	-5.546	63.891	-49.414	1.00	34.57
	ATOM	693	CB	VAL	86	-4.602	65.127	-49.398	1.00	35.39
	ATOM	694	CG1	VAL	86	-3.694	65.079	-48.184	1.00	31.44
	ATOM	695	CG2	VAL	86	-3.769	65.163	-50.664	1.00	38.08
	ATOM	696	C	VAL	86	-6.250	63.833	-48.079	1.00	32.85
40	ATOM	697	O	VAL	86	-7.166	64.609	-47.821	1.00	35.10
	ATOM	698	N	THR	87	-5.842	62.910	-47.226	1.00	33.41
	ATOM	699	CA	THR	87	-6.454	62.856	-45.920	1.00	33.56
	ATOM	700	CB	THR	87	-6.857	61.400	-45.530	1.00	32.70
	ATOM	701	OG1	THR	87	-6.563	61.159	-44.148	1.00	34.29
45	ATOM	702	CG2	THR	87	-6.162	60.397	-46.413	1.00	34.95
	ATOM	703	C	THR	87	-5.528	63.499	-44.879	1.00	30.57
	ATOM	704	O	THR	87	-4.416	63.038	-44.639	1.00	28.08
	ATOM	705	N	LEU	88	-5.994	64.615	-44.326	1.00	28.42
	ATOM	706	CA	LEU	88	-5.282	65.342	-43.291	1.00	31.31
50	ATOM	707	CB	LEU	88	-5.602	66.842	-43.349	1.00	28.21
	ATOM	708	CG	LEU	88	-5.515	67.814	-44.529	1.00	29.12
	ATOM	709	CD1	LEU	88	-4.857	67.185	-45.707	1.00	31.12
	ATOM	710	CD2	LEU	88	-6.904	68.289	-44.879	1.00	28.72
	ATOM	711	C	LEU	88	-5.837	64.814	-41.957	1.00	41.15
55	ATOM	712	O	LEU	88	-7.006	64.430	-41.861	1.00	52.30
	ATOM	713	N	PHE	89	-5.019	64.764	-40.923	1.00	39.23
	ATOM	714	CA	PHE	89	-5.543	64.352	-39.618	1.00	42.27
	ATOM	715	CB	PHE	89	-6.570	65.402	-39.185	1.00	35.63
	ATOM	716	CG	PHE	89	-6.122	66.807	-39.496	1.00	35.06
60	ATOM	717	CD1	PHE	89	-7.004	67.731	-40.047	1.00	31.56
	ATOM	718	CD2	PHE	89	-4.770	67.165	-39.349	1.00	30.96
	ATOM	719	CE1	PHE	89	-6.546	68.985	-40.465	1.00	33.76
	ATOM	720	CE2	PHE	89	-4.305	68.408	-39.763	1.00	27.38
	ATOM	721	CZ	PHE	89	-5.191	69.319	-40.325	1.00	30.29
65	ATOM	722	C	PHE	89	-6.080	62.915	-39.439	1.00	39.99
	ATOM	723	O	PHE	89	-5.314	61.954	-39.566	1.00	45.34

	ATOM	724	N	ARG	90	-7.354	62.734	-39.124	1.00	38.92
	ATOM	725	CA	ARG	90	-7.851	61.366	-38.889	1.00	41.66
	ATOM	726	CB	ARG	90	-7.279	60.384	-39.933	1.00	39.24
	ATOM	727	CG	ARG	90	-6.947	58.979	-39.391	1.00	29.41
5	ATOM	728	CD	ARG	90	-6.408	58.039	-40.486	1.00	31.13
	ATOM	729	NE	ARG	90	-5.838	56.808	-39.932	1.00	31.75
	ATOM	730	CZ	ARG	90	-4.547	56.471	-39.983	1.00	28.86
	ATOM	731	NH1	ARG	90	-3.660	57.255	-40.573	1.00	25.48
	ATOM	732	NH2	ARG	90	-4.133	55.352	-39.412	1.00	29.84
10	ATOM	733	C	ARG	90	-7.529	60.803	-37.492	1.00	39.23
	ATOM	734	O	ARG	90	-6.383	60.837	-37.048	1.00	31.33
	ATOM	735	N	CYS	91	-8.550	60.286	-36.803	1.00	40.12
	ATOM	736	CA	CYS	91	-8.339	59.685	-35.495	1.00	37.09
	ATOM	737	C	CYS	91	-9.035	58.380	-35.236	1.00	37.21
15	ATOM	738	O	CYS	91	-9.920	57.975	-35.980	1.00	37.46
	ATOM	739	CB	CYS	91	-8.674	60.633	-34.367	1.00	40.13
	ATOM	740	SG	CYS	91	-10.350	61.317	-34.117	1.00	41.64
	ATOM	741	N	ILE	92	-8.619	57.729	-34.156	1.00	34.02
	ATOM	742	CA	ILE	92	-9.155	56.429	-33.779	1.00	33.22
20	ATOM	743	CB	ILE	92	-8.148	55.317	-34.137	1.00	32.88
	ATOM	744	CG2	ILE	92	-8.776	53.956	-33.928	1.00	32.61
	ATOM	745	CG1	ILE	92	-7.711	55.467	-35.596	1.00	34.64
	ATOM	746	CD1	ILE	92	-6.572	54.548	-36.009	1.00	32.30
	ATOM	747	C	ILE	92	-9.472	56.324	-32.293	1.00	30.78
25	ATOM	748	O	ILE	92	-8.880	57.007	-31.483	1.00	33.43
	ATOM	749	N	GLN	93	-10.434	55.477	-31.949	1.00	29.35
	ATOM	750	CA	GLN	93	-10.813	55.245	-30.564	1.00	28.69
	ATOM	751	CB	GLN	93	-12.040	56.063	-30.162	1.00	28.17
	ATOM	752	CG	GLN	93	-11.804	57.506	-29.711	1.00	25.87
30	ATOM	753	CD	GLN	93	-10.920	57.629	-28.488	1.00	24.22
	ATOM	754	OE1	GLN	93	-9.707	57.711	-28.602	1.00	28.90
	ATOM	755	NE2	GLN	93	-11.529	57.641	-27.310	1.00	25.29
	ATOM	756	C	GLN	93	-11.169	53.781	-30.439	1.00	31.44
	ATOM	757	O	GLN	93	-11.984	53.282	-31.199	1.00	32.07
35	ATOM	758	N	ASN	94	-10.542	53.076	-29.507	1.00	35.77
	ATOM	759	CA	ASN	94	-10.885	51.682	-29.302	1.00	33.56
	ATOM	760	CB	ASN	94	-9.884	51.002	-28.365	1.00	34.70
	ATOM	761	CG	ASN	94	-8.627	50.566	-29.073	1.00	33.01
	ATOM	762	OD1	ASN	94	-8.684	49.889	-30.086	1.00	30.56
40	ATOM	763	ND2	ASN	94	-7.484	50.936	-28.532	1.00	36.61
	ATOM	764	C	ASN	94	-12.274	51.705	-28.663	1.00	35.80
	ATOM	765	O	ASN	94	-12.583	52.589	-27.860	1.00	36.63
	ATOM	766	N	MET	95	-13.117	50.747	-29.025	1.00	35.73
	ATOM	767	CA	MET	95	-14.468	50.696	-28.480	1.00	34.73
45	ATOM	768	CB	MET	95	-15.481	50.622	-29.626	1.00	32.97
	ATOM	769	CG	MET	95	-15.356	51.724	-30.669	1.00	27.35
	ATOM	770	SD	MET	95	-15.419	53.392	-29.984	1.00	25.14
	ATOM	771	CE	MET	95	-17.115	53.568	-29.544	1.00	15.55
	ATOM	772	C	MET	95	-14.676	49.510	-27.536	1.00	36.36
50	ATOM	773	O	MET	95	-14.005	48.481	-27.662	1.00	40.70
	ATOM	774	N	PRO	96	-15.587	49.651	-26.559	1.00	36.06
	ATOM	775	CD	PRO	96	-16.140	50.923	-26.092	1.00	35.32
	ATOM	776	CA	PRO	96	-15.897	48.594	-25.596	1.00	40.00
	ATOM	777	CB	PRO	96	-16.440	49.354	-24.390	1.00	33.97
55	ATOM	778	CG	PRO	96	-16.024	50.752	-24.617	1.00	37.05
	ATOM	779	C	PRO	96	-16.984	47.723	-26.222	1.00	46.34
	ATOM	780	O	PRO	96	-17.582	48.107	-27.226	1.00	47.59
	ATOM	781	N	GLU	97	-17.256	46.565	-25.629	1.00	52.14
	ATOM	782	CA	GLU	97	-18.279	45.675	-26.164	1.00	57.27
60	ATOM	783	CB	GLU	97	-17.989	44.228	-25.767	1.00	62.29
	ATOM	784	CG	GLU	97	-18.618	43.181	-26.681	1.00	72.31
	ATOM	785	CD	GLU	97	-17.615	42.620	-27.699	1.00	79.34
	ATOM	786	OE1	GLU	97	-16.594	42.001	-27.265	1.00	84.94
	ATOM	787	OE2	GLU	97	-17.844	42.802	-28.927	1.00	81.39
65	ATOM	788	C	GLU	97	-19.637	46.083	-25.603	1.00	56.67
	ATOM	789	O	GLU	97	-20.657	46.013	-26.293	1.00	59.18

	ATOM	790	N	THR	98	-19.638	46.528	-24.351	1.00	57.55
	ATOM	791	CA	THR	98	-20.865	46.913	-23.665	1.00	59.71
	ATOM	792	CB	THR	98	-20.602	47.118	-22.179	1.00	59.24
	ATOM	793	OG1	THR	98	-19.597	48.140	-22.029	1.00	68.81
5	ATOM	794	CG2	THR	98	-20.137	45.796	-21.534	1.00	56.15
	ATOM	795	C	THR	98	-21.594	48.153	-24.180	1.00	60.32
	ATOM	796	O	THR	98	-22.387	48.057	-25.124	1.00	65.03
	ATOM	797	N	LEU	99	-21.349	49.305	-23.553	1.00	54.94
	ATOM	798	CA	LEU	99	-22.020	50.548	-23.945	1.00	51.13
10	ATOM	799	CB	LEU	99	-22.607	51.223	-22.708	1.00	50.56
	ATOM	800	CG	LEU	99	-23.845	50.572	-22.097	1.00	52.80
	ATOM	801	CD1	LEU	99	-24.055	51.070	-20.668	1.00	48.38
	ATOM	802	CD2	LEU	99	-25.051	50.885	-22.969	1.00	50.80
	ATOM	803	C	LEU	99	-21.117	51.538	-24.687	1.00	50.20
15	ATOM	804	O	LEU	99	-20.659	52.535	-24.110	1.00	50.50
	ATOM	805	N	PRO	100	-20.880	51.298	-25.990	1.00	47.97
	ATOM	806	CD	PRO	100	-21.516	50.275	-26.835	1.00	45.68
	ATOM	807	CA	PRO	100	-20.026	52.170	-26.800	1.00	46.54
	ATOM	808	CB	PRO	100	-20.182	51.602	-28.211	1.00	45.50
20	ATOM	809	CG	PRO	100	-20.542	50.172	-27.973	1.00	45.80
	ATOM	810	C	PRO	100	-20.435	53.631	-26.732	1.00	46.78
	ATOM	811	O	PRO	100	-21.568	53.981	-27.038	1.00	52.13
	ATOM	812	N	ASN	101	-19.507	54.481	-26.327	1.00	42.71
	ATOM	813	CA	ASN	101	-19.751	55.908	-26.245	1.00	41.37
25	ATOM	814	CB	ASN	101	-20.619	56.231	-25.035	1.00	43.24
	ATOM	815	CG	ASN	101	-22.098	56.062	-25.315	1.00	45.93
	ATOM	816	OD1	ASN	101	-22.693	56.837	-26.067	1.00	45.33
	ATOM	817	ND2	ASN	101	-22.704	55.040	-24.714	1.00	48.11
	ATOM	818	C	ASN	101	-18.412	56.608	-26.128	1.00	42.66
30	ATOM	819	O	ASN	101	-17.972	56.950	-25.028	1.00	44.79
	ATOM	820	N	ASN	102	-17.745	56.820	-27.258	1.00	40.46
	ATOM	821	CA	ASN	102	-16.462	57.478	-27.194	1.00	37.90
	ATOM	822	CB	ASN	102	-15.374	56.584	-27.769	1.00	34.01
	ATOM	823	CG	ASN	102	-14.803	55.642	-26.727	1.00	32.44
35	ATOM	824	OD1	ASN	102	-14.595	56.030	-25.581	1.00	29.07
	ATOM	825	ND2	ASN	102	-14.541	54.405	-27.121	1.00	33.79
	ATOM	826	C	ASN	102	-16.323	58.884	-27.746	1.00	39.20
	ATOM	827	O	ASN	102	-16.054	59.807	-26.987	1.00	46.07
	ATOM	828	N	SER	103	-16.501	59.087	-29.037	1.00	35.42
40	ATOM	829	CA	SER	103	-16.322	60.450	-29.565	1.00	40.28
	ATOM	830	CB	SER	103	-17.177	61.485	-28.793	1.00	40.68
	ATOM	831	OG	SER	103	-16.417	62.246	-27.862	1.00	31.76
	ATOM	832	C	SER	103	-14.831	60.863	-29.516	1.00	35.24
	ATOM	833	O	SER	103	-14.189	60.861	-28.467	1.00	24.65
45	ATOM	834	N	CYS	104	-14.293	61.185	-30.684	1.00	32.88
	ATOM	835	CA	CYS	104	-12.915	61.584	-30.808	1.00	31.77
	ATOM	836	C	CYS	104	-12.875	62.942	-31.531	1.00	30.58
	ATOM	837	O	CYS	104	-13.580	63.156	-32.513	1.00	29.23
	ATOM	838	CB	CYS	104	-12.103	60.511	-31.585	1.00	28.43
50	ATOM	839	SG	CYS	104	-10.518	61.232	-32.039	1.00	46.07
	ATOM	840	N	TYR	105	-12.071	63.869	-31.014	1.00	30.34
	ATOM	841	CA	TYR	105	-11.900	65.190	-31.617	1.00	26.12
	ATOM	842	CB	TYR	105	-12.072	66.287	-30.567	1.00	27.03
	ATOM	843	CG	TYR	105	-11.726	67.688	-31.045	1.00	25.61
55	ATOM	844	CD1	TYR	105	-12.718	68.575	-31.477	1.00	24.90
	ATOM	845	CE1	TYR	105	-12.402	69.853	-31.914	1.00	26.08
	ATOM	846	CD2	TYR	105	-10.406	68.129	-31.069	1.00	25.41
	ATOM	847	CE2	TYR	105	-10.083	69.406	-31.510	1.00	28.91
	ATOM	848	CZ	TYR	105	-11.085	70.263	-31.929	1.00	27.58
60	ATOM	849	OH	TYR	105	-10.762	71.533	-32.352	1.00	29.89
	ATOM	850	C	TYR	105	-10.489	65.273	-32.192	1.00	29.20
	ATOM	851	O	TYR	105	-9.545	64.730	-31.626	1.00	33.10
	ATOM	852	N	SER	106	-10.338	65.950	-33.319	1.00	30.48
	ATOM	853	CA	SER	106	-9.024	66.086	-33.925	1.00	29.22
65	ATOM	854	CB	SER	106	-8.699	64.844	-34.759	1.00	28.30
	ATOM	855	OG	SER	106	-7.405	64.918	-35.324	1.00	28.62

	ATOM	856	C	SER	106	-9.037	67.327	-34.801	1.00	29.94
	ATOM	857	O	SER	106	-10.047	67.624	-35.427	1.00	29.72
	ATOM	858	N	ALA	107	-7.927	68.060	-34.825	1.00	27.64
	ATOM	859	CA	ALA	107	-7.829	69.269	-35.632	1.00	24.33
5	ATOM	860	CB	ALA	107	-8.412	70.451	-34.878	1.00	21.30
	ATOM	861	C	ALA	107	-6.393	69.568	-36.032	1.00	25.56
	ATOM	862	O	ALA	107	-5.451	69.009	-35.485	1.00	28.03
	ATOM	863	N	GLY	108	-6.233	70.460	-36.995	1.00	26.07
	ATOM	864	CA	GLY	108	-4.908	70.832	-37.446	1.00	25.49
10	ATOM	865	C	GLY	108	-4.994	71.929	-38.481	1.00	29.03
	ATOM	866	O	GLY	108	-6.085	72.321	-38.872	1.00	32.20
	ATOM	867	N	ILE	109	-3.850	72.427	-38.929	1.00	29.34
	ATOM	868	CA	ILE	109	-3.817	73.486	-39.926	1.00	27.71
	ATOM	869	CB	ILE	109	-2.853	74.607	-39.505	1.00	27.48
15	ATOM	870	CG2	ILE	109	-2.812	75.690	-40.562	1.00	26.46
	ATOM	871	CG1	ILE	109	-3.303	75.187	-38.165	1.00	26.16
	ATOM	872	CD1	ILE	109	-2.341	76.178	-37.573	1.00	29.07
	ATOM	873	C	ILE	109	-3.367	72.920	-41.259	1.00	28.66
	ATOM	874	O	ILE	109	-2.505	72.044	-41.311	1.00	30.59
20	ATOM	875	N	ALA	110	-3.953	73.413	-42.341	1.00	25.60
	ATOM	876	CA	ALA	110	-3.600	72.954	-43.679	1.00	26.36
	ATOM	877	CB	ALA	110	-4.488	71.800	-44.081	1.00	20.26
	ATOM	878	C	ALA	110	-3.758	74.090	-44.674	1.00	29.61
	ATOM	879	O	ALA	110	-4.525	75.014	-44.447	1.00	27.61
25	ATOM	880	N	LYS	111	-3.018	74.039	-45.771	1.00	33.77
	ATOM	881	CA	LYS	111	-3.153	75.074	-46.770	1.00	35.09
	ATOM	882	CB	LYS	111	-1.814	75.413	-47.412	1.00	39.38
	ATOM	883	CG	LYS	111	-1.997	76.383	-48.572	1.00	49.74
	ATOM	884	CD	LYS	111	-0.858	77.355	-48.718	1.00	53.95
30	ATOM	885	CE	LYS	111	-1.193	78.394	-49.778	1.00	54.28
	ATOM	886	NZ	LYS	111	-0.055	79.359	-49.958	1.00	60.85
	ATOM	887	C	LYS	111	-4.133	74.582	-47.828	1.00	35.28
	ATOM	888	O	LYS	111	-3.969	73.502	-48.384	1.00	36.58
	ATOM	889	N	LEU	112	-5.158	75.383	-48.089	1.00	32.32
35	ATOM	890	CA	LEU	112	-6.180	75.036	-49.057	1.00	29.67
	ATOM	891	CB	LEU	112	-7.539	74.948	-48.367	1.00	27.04
	ATOM	892	CG	LEU	112	-7.597	74.101	-47.097	1.00	24.38
	ATOM	893	CD1	LEU	112	-8.966	74.226	-46.459	1.00	27.56
	ATOM	894	CD2	LEU	112	-7.289	72.666	-47.425	1.00	19.52
40	ATOM	895	C	LEU	112	-6.237	76.085	-50.158	1.00	32.58
	ATOM	896	O	LEU	112	-5.692	77.178	-50.013	1.00	31.51
	ATOM	897	N	GLU	113	-6.906	75.751	-51.255	1.00	35.75
	ATOM	898	CA	GLU	113	-7.028	76.666	-52.376	1.00	40.08
	ATOM	899	CB	GLU	113	-6.182	76.205	-53.548	1.00	44.23
45	ATOM	900	CG	GLU	113	-4.774	75.806	-53.219	1.00	50.12
	ATOM	901	CD	GLU	113	-3.901	75.872	-54.451	1.00	58.08
	ATOM	902	OE1	GLU	113	-4.397	75.505	-55.557	1.00	61.13
	ATOM	903	OE2	GLU	113	-2.723	76.292	-54.311	1.00	63.64
	ATOM	904	C	GLU	113	-8.450	76.764	-52.871	1.00	41.63
50	ATOM	905	O	GLU	113	-9.265	75.887	-52.618	1.00	48.29
	ATOM	906	N	GLU	114	-8.729	77.843	-53.596	1.00	40.97
	ATOM	907	CA	GLU	114	-10.039	78.073	-54.191	1.00	36.68
	ATOM	908	CB	GLU	114	-9.953	79.164	-55.238	1.00	41.19
	ATOM	909	CG	GLU	114	-10.324	80.536	-54.802	1.00	46.58
55	ATOM	910	CD	GLU	114	-10.810	81.348	-55.984	1.00	51.12
	ATOM	911	OE1	GLU	114	-11.886	81.001	-56.530	1.00	51.88
	ATOM	912	OE2	GLU	114	-10.111	82.315	-56.374	1.00	57.08
	ATOM	913	C	GLU	114	-10.469	76.818	-54.922	1.00	35.51
	ATOM	914	O	GLU	114	-9.766	76.353	-55.812	1.00	38.92
60	ATOM	915	N	GLY	115	-11.623	76.276	-54.566	1.00	32.99
	ATOM	916	CA	GLY	115	-12.083	75.092	-55.258	1.00	31.79
	ATOM	917	C	GLY	115	-11.864	73.812	-54.498	1.00	34.39
	ATOM	918	O	GLY	115	-12.468	72.800	-54.825	1.00	39.20
	ATOM	919	N	ASP	116	-10.984	73.839	-53.506	1.00	35.66
65	ATOM	920	CA	ASP	116	-10.742	72.652	-52.705	1.00	38.09
	ATOM	921	CB	ASP	116	-9.575	72.863	-51.732	1.00	36.00

	ATOM	922	CG	ASP	116	-8.222	72.777	-52.403	1.00	35.09
	ATOM	923	OD1	ASP	116	-8.136	72.273	-53.542	1.00	33.81
	ATOM	924	OD2	ASP	116	-7.233	73.199	-51.776	1.00	35.24
	ATOM	925	C	ASP	116	-12.005	72.378	-51.903	1.00	39.38
5	ATOM	926	O	ASP	116	-12.777	73.296	-51.609	1.00	37.91
	ATOM	927	N	GLU	117	-12.228	71.116	-51.560	1.00	41.54
	ATOM	928	CA	GLU	117	-13.388	70.757	-50.756	1.00	37.92
	ATOM	929	CB	GLU	117	-14.377	69.926	-51.565	1.00	38.90
	ATOM	930	CG	GLU	117	-14.876	70.609	-52.812	1.00	48.15
10	ATOM	931	CD	GLU	117	-15.886	69.763	-53.560	1.00	52.62
	ATOM	932	OE1	GLU	117	-15.712	68.514	-53.604	1.00	55.49
	ATOM	933	OE2	GLU	117	-16.848	70.354	-54.110	1.00	57.10
	ATOM	934	C	GLU	117	-12.905	69.947	-49.570	1.00	34.02
	ATOM	935	O	GLU	117	-11.949	69.181	-49.686	1.00	33.31
15	ATOM	936	N	LEU	118	-13.545	70.139	-48.425	1.00	30.68
	ATOM	937	CA	LEU	118	-13.199	69.399	-47.223	1.00	27.14
	ATOM	938	CB	LEU	118	-12.979	70.353	-46.049	1.00	26.49
	ATOM	939	CG	LEU	118	-11.801	71.319	-46.109	1.00	26.21
	ATOM	940	CD1	LEU	118	-11.828	72.217	-44.894	1.00	23.05
20	ATOM	941	CD2	LEU	118	-10.500	70.546	-46.172	1.00	25.38
	ATOM	942	C	LEU	118	-14.354	68.464	-46.889	1.00	28.33
	ATOM	943	O	LEU	118	-15.517	68.803	-47.115	1.00	31.46
	ATOM	944	N	GLN	119	-14.044	67.283	-46.364	1.00	28.96
	ATOM	945	CA	GLN	119	-15.093	66.348	-45.976	1.00	29.13
25	ATOM	946	CB	GLN	119	-15.480	65.469	-47.156	1.00	28.23
	ATOM	947	CG	GLN	119	-14.428	64.479	-47.554	1.00	36.38
	ATOM	948	CD	GLN	119	-14.828	63.677	-48.769	1.00	36.57
	ATOM	949	OE1	GLN	119	-14.229	62.642	-49.059	1.00	40.70
	ATOM	950	NE2	GLN	119	-15.834	64.152	-49.493	1.00	36.80
30	ATOM	951	C	GLN	119	-14.678	65.486	-44.788	1.00	29.43
	ATOM	952	O	GLN	119	-13.494	65.256	-44.557	1.00	28.80
	ATOM	953	N	LEU	120	-15.674	65.035	-44.033	1.00	32.26
	ATOM	954	CA	LEU	120	-15.481	64.197	-42.852	1.00	29.96
	ATOM	955	CB	LEU	120	-16.298	64.767	-41.683	1.00	31.17
35	ATOM	956	CG	LEU	120	-16.158	64.338	-40.217	1.00	30.26
	ATOM	957	CD1	LEU	120	-16.135	62.835	-40.087	1.00	27.10
	ATOM	958	CD2	LEU	120	-14.903	64.943	-39.654	1.00	30.96
	ATOM	959	C	LEU	120	-15.991	62.802	-43.216	1.00	30.51
	ATOM	960	O	LEU	120	-17.187	62.604	-43.440	1.00	32.88
40	ATOM	961	N	ALA	121	-15.083	61.834	-43.268	1.00	27.40
	ATOM	962	CA	ALA	121	-15.448	60.470	-43.626	1.00	25.55
	ATOM	963	CB	ALA	121	-14.751	60.086	-44.922	1.00	19.41
	ATOM	964	C	ALA	121	-15.139	59.430	-42.549	1.00	29.89
	ATOM	965	O	ALA	121	-14.091	59.474	-41.907	1.00	35.88
45	ATOM	966	N	ILE	122	-16.069	58.500	-42.355	1.00	30.33
	ATOM	967	CA	ILE	122	-15.895	57.424	-41.388	1.00	30.27
	ATOM	968	CB	ILE	122	-17.136	57.251	-40.507	1.00	29.27
	ATOM	969	CG2	ILE	122	-16.868	56.188	-39.466	1.00	30.38
	ATOM	970	CG1	ILE	122	-17.484	58.580	-39.829	1.00	26.03
50	ATOM	971	CD1	ILE	122	-18.721	58.539	-38.952	1.00	29.21
	ATOM	972	C	ILE	122	-15.680	56.157	-42.210	1.00	33.20
	ATOM	973	O	ILE	122	-16.576	55.719	-42.928	1.00	33.30
	ATOM	974	N	PRO	123	-14.481	55.556	-42.118	1.00	34.70
	ATOM	975	CD	PRO	123	-13.349	56.036	-41.319	1.00	34.40
55	ATOM	976	CA	PRO	123	-14.104	54.337	-42.846	1.00	36.49
	ATOM	977	CB	PRO	123	-12.585	54.271	-42.667	1.00	30.22
	ATOM	978	CG	PRO	123	-12.201	55.641	-42.186	1.00	36.72
	ATOM	979	C	PRO	123	-14.770	53.064	-42.326	1.00	39.64
	ATOM	980	O	PRO	123	-14.078	52.109	-41.959	1.00	40.48
60	ATOM	981	N	ARG	124	-16.099	53.049	-42.298	1.00	44.58
	ATOM	982	CA	ARG	124	-16.841	51.890	-41.822	1.00	48.04
	ATOM	983	CB	ARG	124	-17.065	51.986	-40.317	1.00	52.91
	ATOM	984	CG	ARG	124	-15.788	52.173	-39.518	1.00	60.50
	ATOM	985	CD	ARG	124	-15.489	50.896	-38.764	1.00	72.57
65	ATOM	986	NE	ARG	124	-14.177	50.944	-38.094	1.00	84.90
	ATOM	987	CZ	ARG	124	-13.020	50.532	-38.627	1.00	85.44

	ATOM	988	NH1	ARG	124	-12.992	50.030	-39.863	1.00	87.69
	ATOM	989	NH2	ARG	124	-11.886	50.626	-37.922	1.00	82.78
	ATOM	990	C	ARG	124	-18.177	51.745	-42.532	1.00	48.39
	ATOM	991	O	ARG	124	-18.799	52.739	-42.909	1.00	45.26
5	ATOM	992	N	GLU	125	-18.602	50.495	-42.696	1.00	52.84
	ATOM	993	CA	GLU	125	-19.848	50.149	-43.377	1.00	55.02
	ATOM	994	CB	GLU	125	-20.113	48.648	-43.229	1.00	63.62
	ATOM	995	CG	GLU	125	-19.544	48.038	-41.945	1.00	70.62
	ATOM	996	CD	GLU	125	-18.051	48.298	-41.786	1.00	74.76
10	ATOM	997	OE1	GLU	125	-17.258	47.781	-42.619	1.00	75.75
	ATOM	998	OE2	GLU	125	-17.673	49.018	-40.829	1.00	80.71
	ATOM	999	C	GLU	125	-21.073	50.936	-42.931	1.00	53.44
	ATOM	1000	O	GLU	125	-21.588	51.758	-43.690	1.00	57.50
	ATOM	1001	N	ASN	126	-21.575	50.668	-41.734	1.00	47.18
15	ATOM	1002	CA	ASN	126	-22.724	51.420	-41.249	1.00	47.59
	ATOM	1003	CB	ASN	126	-23.995	50.575	-41.244	1.00	51.03
	ATOM	1004	CG	ASN	126	-24.898	50.882	-42.429	1.00	57.07
	ATOM	1005	OD1	ASN	126	-24.631	50.467	-43.567	1.00	61.15
	ATOM	1006	ND2	ASN	126	-25.968	51.635	-42.172	1.00	56.63
20	ATOM	1007	C	ASN	126	-22.406	51.893	-39.857	1.00	47.06
	ATOM	1008	O	ASN	126	-22.971	51.422	-38.866	1.00	45.61
	ATOM	1009	N	ALA	127	-21.474	52.835	-39.803	1.00	43.50
	ATOM	1010	CA	ALA	127	-21.009	53.394	-38.554	1.00	38.26
	ATOM	1011	CB	ALA	127	-20.169	54.619	-38.832	1.00	37.97
25	ATOM	1012	C	ALA	127	-22.148	53.748	-37.618	1.00	37.16
	ATOM	1013	O	ALA	127	-23.092	54.437	-38.000	1.00	33.14
	ATOM	1014	N	GLN	128	-22.062	53.247	-36.392	1.00	37.87
	ATOM	1015	CA	GLN	128	-23.058	53.562	-35.379	1.00	37.19
	ATOM	1016	CB	GLN	128	-23.147	52.423	-34.368	1.00	40.23
30	ATOM	1017	CG	GLN	128	-23.737	51.172	-34.973	1.00	38.14
	ATOM	1018	CD	GLN	128	-25.075	51.454	-35.615	1.00	39.51
	ATOM	1019	OE1	GLN	128	-26.057	51.756	-34.929	1.00	40.36
	ATOM	1020	NE2	GLN	128	-25.121	51.383	-36.945	1.00	41.53
	ATOM	1021	C	GLN	128	-22.559	54.854	-34.738	1.00	35.55
35	ATOM	1022	O	GLN	128	-21.662	54.858	-33.890	1.00	35.57
	ATOM	1023	N	ILE	129	-23.156	55.951	-35.178	1.00	34.64
	ATOM	1024	CA	ILE	129	-22.772	57.295	-34.768	1.00	29.60
	ATOM	1025	CB	ILE	129	-22.211	58.031	-36.039	1.00	31.03
	ATOM	1026	CG2	ILE	129	-22.787	59.414	-36.201	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-20.700	58.037	-35.989	1.00	26.37
	ATOM	1028	CD1	ILE	129	-20.123	56.671	-35.985	1.00	34.70
	ATOM	1029	C	ILE	129	-23.922	58.083	-34.150	1.00	29.55
	ATOM	1030	O	ILE	129	-25.082	57.814	-34.425	1.00	33.42
	ATOM	1031	N	SER	130	-23.601	59.054	-33.303	1.00	27.67
45	ATOM	1032	CA	SER	130	-24.633	59.899	-32.709	1.00	25.00
	ATOM	1033	CB	SER	130	-24.240	60.348	-31.312	1.00	22.09
	ATOM	1034	OG	SER	130	-25.138	61.350	-30.855	1.00	20.89
	ATOM	1035	C	SER	130	-24.784	61.129	-33.589	1.00	26.03
	ATOM	1036	O	SER	130	-23.795	61.702	-34.011	1.00	30.77
50	ATOM	1037	N	LEU	131	-26.012	61.544	-33.863	1.00	24.49
	ATOM	1038	CA	LEU	131	-26.231	62.704	-34.713	1.00	28.07
	ATOM	1039	CB	LEU	131	-27.226	62.366	-35.824	1.00	28.87
	ATOM	1040	CG	LEU	131	-26.641	61.780	-37.108	1.00	30.76
	ATOM	1041	CD1	LEU	131	-25.674	60.673	-36.806	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-27.770	61.276	-37.968	1.00	33.37
	ATOM	1043	C	LEU	131	-26.693	63.952	-33.980	1.00	33.66
	ATOM	1044	O	LEU	131	-27.420	64.772	-34.541	1.00	37.67
	ATOM	1045	N	ASP	132	-26.273	64.096	-32.728	1.00	34.97
	ATOM	1046	CA	ASP	132	-26.628	65.274	-31.942	1.00	34.31
60	ATOM	1047	CB	ASP	132	-26.556	64.973	-30.445	1.00	42.68
	ATOM	1048	CG	ASP	132	-27.743	64.162	-29.953	1.00	47.98
	ATOM	1049	OD1	ASP	132	-27.693	63.676	-28.796	1.00	50.41
	ATOM	1050	OD2	ASP	132	-28.727	64.022	-30.723	1.00	46.42
	ATOM	1051	C	ASP	132	-25.662	66.392	-32.295	1.00	34.32
65	ATOM	1052	O	ASP	132	-24.454	66.188	-32.349	1.00	32.87
	ATOM	1053	N	GLY	133	-26.210	67.575	-32.537	1.00	34.53

	ATOM	1054	CA	GLY	133	-25.396	68.713	-32.915	1.00	36.93
	ATOM	1055	C	GLY	133	-24.245	69.079	-31.999	1.00	35.06
	ATOM	1056	O	GLY	133	-23.295	69.733	-32.429	1.00	40.44
	ATOM	1057	N	ASP	134	-24.308	68.664	-30.742	1.00	31.08
5	ATOM	1058	CA	ASP	134	-23.248	69.009	-29.816	1.00	27.63
	ATOM	1059	CB	ASP	134	-23.834	69.309	-28.434	1.00	28.43
	ATOM	1060	CG	ASP	134	-24.580	68.135	-27.839	1.00	31.78
	ATOM	1061	OD1	ASP	134	-25.201	67.370	-28.601	1.00	35.68
	ATOM	1062	OD2	ASP	134	-24.562	67.985	-26.599	1.00	32.53
10	ATOM	1063	C	ASP	134	-22.136	67.980	-29.729	1.00	25.08
	ATOM	1064	O	ASP	134	-21.039	68.302	-29.331	1.00	19.71
	ATOM	1065	N	VAL	135	-22.393	66.747	-30.136	1.00	26.84
	ATOM	1066	CA	VAL	135	-21.345	65.740	-30.059	1.00	23.72
	ATOM	1067	CB	VAL	135	-21.879	64.421	-29.496	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-22.189	64.602	-28.035	1.00	22.76
	ATOM	1069	CG2	VAL	135	-23.112	63.984	-30.256	1.00	22.12
	ATOM	1070	C	VAL	135	-20.596	65.473	-31.361	1.00	26.54
	ATOM	1071	O	VAL	135	-19.458	65.025	-31.325	1.00	29.26
	ATOM	1072	N	THR	136	-21.217	65.729	-32.511	1.00	26.39
20	ATOM	1073	CA	THR	136	-20.503	65.531	-33.771	1.00	28.40
	ATOM	1074	CB	THR	136	-20.934	64.210	-34.492	1.00	28.21
	ATOM	1075	OG1	THR	136	-21.812	64.502	-35.577	1.00	37.13
	ATOM	1076	CG2	THR	136	-21.622	63.275	-33.529	1.00	26.90
	ATOM	1077	C	THR	136	-20.666	66.757	-34.682	1.00	31.60
25	ATOM	1078	O	THR	136	-21.773	67.111	-35.096	1.00	27.29
	ATOM	1079	N	PHE	137	-19.538	67.411	-34.965	1.00	31.45
	ATOM	1080	CA	PHE	137	-19.519	68.622	-35.777	1.00	26.64
	ATOM	1081	CB	PHE	137	-19.619	69.828	-34.853	1.00	26.84
	ATOM	1082	CG	PHE	137	-18.763	69.722	-33.625	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-17.430	70.119	-33.651	1.00	27.77
	ATOM	1084	CD2	PHE	137	-19.288	69.235	-32.440	1.00	23.70
	ATOM	1085	CE1	PHE	137	-16.638	70.036	-32.517	1.00	27.32
	ATOM	1086	CE2	PHE	137	-18.501	69.147	-31.302	1.00	26.31
	ATOM	1087	CZ	PHE	137	-17.174	69.550	-31.343	1.00	29.43
35	ATOM	1088	C	PHE	137	-18.284	68.710	-36.670	1.00	28.06
	ATOM	1089	O	PHE	137	-17.325	67.979	-36.470	1.00	25.04
	ATOM	1090	N	PHE	138	-18.314	69.614	-37.649	1.00	29.81
	ATOM	1091	CA	PHE	138	-17.222	69.749	-38.610	1.00	29.93
	ATOM	1092	CB	PHE	138	-17.703	69.172	-39.940	1.00	27.47
40	ATOM	1093	CG	PHE	138	-16.625	68.966	-40.949	1.00	28.49
	ATOM	1094	CD1	PHE	138	-15.299	68.812	-40.561	1.00	27.65
	ATOM	1095	CD2	PHE	138	-16.941	68.921	-42.305	1.00	31.06
	ATOM	1096	CE1	PHE	138	-14.304	68.618	-41.510	1.00	30.90
	ATOM	1097	CE2	PHE	138	-15.955	68.728	-43.265	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-14.634	68.577	-42.868	1.00	27.41
	ATOM	1099	C	PHE	138	-16.692	71.189	-38.761	1.00	30.91
	ATOM	1100	O	PHE	138	-17.443	72.127	-39.001	1.00	24.94
	ATOM	1101	N	GLY	139	-15.366	71.302	-38.642	1.00	36.94
	ATOM	1102	CA	GLY	139	-14.609	72.553	-38.646	1.00	36.08
50	ATOM	1103	C	GLY	139	-14.471	73.610	-39.714	1.00	36.48
	ATOM	1104	O	GLY	139	-15.461	74.071	-40.251	1.00	43.61
	ATOM	1105	N	ALA	140	-13.228	74.035	-39.950	1.00	36.60
	ATOM	1106	CA	ALA	140	-12.856	75.066	-40.938	1.00	34.94
	ATOM	1107	CB	ALA	140	-13.628	74.860	-42.229	1.00	36.94
55	ATOM	1108	C	ALA	140	-12.948	76.546	-40.519	1.00	32.53
	ATOM	1109	O	ALA	140	-14.025	77.128	-40.449	1.00	27.96
	ATOM	1110	N	LEU	141	-11.786	77.146	-40.271	1.00	33.23
	ATOM	1111	CA	LEU	141	-11.660	78.556	-39.877	1.00	36.11
	ATOM	1112	CB	LEU	141	-11.457	78.652	-38.365	1.00	37.53
60	ATOM	1113	CG	LEU	141	-11.205	80.002	-37.685	1.00	38.87
	ATOM	1114	CD1	LEU	141	-11.307	79.816	-36.190	1.00	40.67
	ATOM	1115	CD2	LEU	141	-9.836	80.552	-38.031	1.00	37.53
	ATOM	1116	C	LEU	141	-10.447	79.151	-40.600	1.00	38.89
	ATOM	1117	O	LEU	141	-9.357	78.585	-40.555	1.00	39.25
65	ATOM	1118	N	LYS	142	-10.616	80.291	-41.261	1.00	42.86
	ATOM	1119	CA	LYS	142	-9.489	80.878	-41.977	1.00	42.96

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	ATOM	1120	CB	LYS	142	-9.984	81.705	-43.160	1.00	41.80
	ATOM	1121	CG	LYS	142	-8.850	82.356	-43.916	1.00	46.88
	ATOM	1122	CD	LYS	142	-9.243	82.769	-45.319	1.00	49.58
	ATOM	1123	CE	LYS	142	-8.052	83.409	-46.012	1.00	51.07
5	ATOM	1124	NZ	LYS	142	-8.339	83.765	-47.423	1.00	56.20
	ATOM	1125	C	LYS	142	-8.550	81.723	-41.123	1.00	43.41
	ATOM	1126	O	LYS	142	-8.985	82.612	-40.397	1.00	44.98
	ATOM	1127	N	LEU	143	-7.254	81.439	-41.220	1.00	42.81
	ATOM	1128	CA	LEU	143	-6.244	82.173	-40.462	1.00	40.65
10	ATOM	1129	CB	LEU	143	-4.999	81.311	-40.264	1.00	36.56
	ATOM	1130	CG	LEU	143	-5.155	79.956	-39.572	1.00	36.17
	ATOM	1131	CD1	LEU	143	-3.809	79.255	-39.542	1.00	37.95
	ATOM	1132	CD2	LEU	143	-5.688	80.148	-38.169	1.00	27.38
	ATOM	1133	C	LEU	143	-5.839	83.442	-41.204	1.00	43.14
15	ATOM	1134	O	LEU	143	-5.855	83.481	-42.439	1.00	46.28
	ATOM	1135	N	LEU	144	-5.475	84.478	-40.460	1.00	42.63
	ATOM	1136	CA	LEU	144	-5.055	85.724	-41.084	1.00	44.11
	ATOM	1137	CB	LEU	144	-5.159	86.881	-40.095	1.00	44.88
	ATOM	1138	CG	LEU	144	-6.583	87.205	-39.649	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-6.562	88.287	-38.604	1.00	47.71
	ATOM	1140	CD2	LEU	144	-7.408	87.642	-40.846	1.00	47.87
	ATOM	1141	C	LEU	144	-3.621	85.602	-41.561	1.00	45.66
	ATOM	1142	O	LEU	144	-2.903	84.720	-41.055	1.00	46.37
	ATOM	1143	OXT	LEU	144	-3.230	86.404	-42.430	1.00	50.66
25	END					31.712	6.654	-112.989	0.00	0.00

TABLE 4

	11								
	ATOM	1	CB	VAL	1	-0.474	91.822	-41.232	1.00 69.28
5	ATOM	2	CG1	VAL	1	-1.417	91.974	-42.446	1.00 68.26
	ATOM	3	CG2	VAL	1	0.238	93.141	-40.914	1.00 71.75
	ATOM	4	C	VAL	1	-1.632	89.844	-40.205	1.00 64.86
	ATOM	5	O	VAL	1	-2.786	89.510	-40.512	1.00 64.02
	ATOM	6	N	VAL	1	-0.491	91.505	-38.726	1.00 65.10
10	ATOM	7	CA	VAL	1	-1.277	91.328	-39.985	1.00 65.98
	ATOM	8	N	THR	2	-0.640	88.963	-40.058	1.00 61.42
	ATOM	9	CA	THR	2	-0.864	87.529	-40.238	1.00 56.64
	ATOM	10	CB	THR	2	0.008	86.947	-41.365	1.00 56.77
	ATOM	11	OG1	THR	2	1.380	87.248	-41.098	1.00 57.45
15	ATOM	12	CG2	THR	2	-0.388	87.526	-42.718	1.00 55.68
	ATOM	13	C	THR	2	-0.538	86.749	-38.979	1.00 53.08
	ATOM	14	O	THR	2	0.046	87.278	-38.038	1.00 52.00
	ATOM	15	N	GLN	3	-0.906	85.474	-38.978	1.00 50.34
	ATOM	16	CA	GLN	3	-0.654	84.607	-37.834	1.00 45.97
20	ATOM	17	CB	GLN	3	-1.947	83.927	-37.399	1.00 46.09
	ATOM	18	CG	GLN	3	-3.112	84.869	-37.247	1.00 49.60
	ATOM	19	CD	GLN	3	-4.344	84.166	-36.739	1.00 50.43
	ATOM	20	OE1	GLN	3	-4.377	83.716	-35.598	1.00 50.94
	ATOM	21	NE2	GLN	3	-5.365	84.058	-37.586	1.00 49.99
25	ATOM	22	C	GLN	3	0.377	83.547	-38.174	1.00 42.41
	ATOM	23	O	GLN	3	0.114	82.629	-38.945	1.00 37.34
	ATOM	24	N	ASP	4	1.559	83.680	-37.595	1.00 40.32
	ATOM	25	CA	ASP	4	2.622	82.723	-37.843	1.00 39.37
	ATOM	26	CB	ASP	4	3.935	83.223	-37.235	1.00 44.56
30	ATOM	27	CG	ASP	4	4.443	84.482	-37.904	1.00 45.76
	ATOM	28	OD1	ASP	4	3.630	85.164	-38.569	1.00 48.58
	ATOM	29	OD2	ASP	4	5.645	84.791	-37.760	1.00 45.55
	ATOM	30	C	ASP	4	2.248	81.396	-37.220	1.00 36.30
	ATOM	31	O	ASP	4	1.583	81.348	-36.188	1.00 36.29
35	ATOM	32	N	CYS	5	2.667	80.316	-37.860	1.00 35.11
	ATOM	33	CA	CYS	5	2.405	78.984	-37.346	1.00 35.61
	ATOM	34	CB	CYS	5	0.955	78.567	-37.605	1.00 34.96
	ATOM	35	SG	CYS	5	0.320	78.978	-39.217	1.00 35.34
	ATOM	36	C	CYS	5	3.370	77.998	-37.971	1.00 31.80
40	ATOM	37	O	CYS	5	3.849	78.203	-39.080	1.00 31.85
	ATOM	38	N	LEU	6	3.680	76.947	-37.230	1.00 27.72
	ATOM	39	CA	LEU	6	4.581	75.914	-37.698	1.00 30.80
	ATOM	40	CB	LEU	6	5.965	76.107	-37.090	1.00 29.25
	ATOM	41	CG	LEU	6	7.035	75.062	-37.412	1.00 29.47
45	ATOM	42	CD1	LEU	6	8.402	75.678	-37.281	1.00 30.65
	ATOM	43	CD2	LEU	6	6.908	73.899	-36.475	1.00 35.27
	ATOM	44	C	LEU	6	3.995	74.583	-37.271	1.00 31.83
	ATOM	45	O	LEU	6	3.672	74.396	-36.104	1.00 33.45
	ATOM	46	N	GLN	7	3.844	73.661	-38.215	1.00 32.23
50	ATOM	47	CA	GLN	7	3.278	72.366	-37.906	1.00 30.26
	ATOM	48	CB	GLN	7	1.998	72.162	-38.706	1.00 29.56
	ATOM	49	CG	GLN	7	1.151	70.988	-38.267	1.00 25.09
	ATOM	50	CD	GLN	7	-0.214	71.003	-38.927	1.00 26.50
	ATOM	51	OE1	GLN	7	-0.346	70.758	-40.126	1.00 24.40
55	ATOM	52	NE2	GLN	7	-1.235	71.310	-38.148	1.00 27.03
	ATOM	53	C	GLN	7	4.275	71.270	-38.216	1.00 32.11
	ATOM	54	O	GLN	7	4.995	71.332	-39.205	1.00 34.24
	ATOM	55	N	LEU	8	4.317	70.275	-37.343	1.00 33.73
	ATOM	56	CA	LEU	8	5.216	69.141	-37.489	1.00 35.41
60	ATOM	57	CB	LEU	8	6.115	69.021	-36.247	1.00 32.44
	ATOM	58	CG	LEU	8	7.500	69.684	-36.214	1.00 27.69
	ATOM	59	CD1	LEU	8	7.630	70.760	-37.261	1.00 26.43
	ATOM	60	CD2	LEU	8	7.734	70.228	-34.831	1.00 18.64
	ATOM	61	C	LEU	8	4.416	67.863	-37.687	1.00 36.87
65	ATOM	62	O	LEU	8	3.318	67.704	-37.156	1.00 36.05
	ATOM	63	N	ILE	9	4.989	66.948	-38.454	1.00 39.25

	ATOM	64	CA	ILE	9	4.358	65.677	-38.769	1.00	35.57
	ATOM	65	CB	ILE	9	4.007	65.658	-40.278	1.00	34.28
	ATOM	66	CG2	ILE	9	4.409	64.359	-40.930	1.00	38.20
	ATOM	67	CG1	ILE	9	2.529	65.915	-40.455	1.00	35.22
5	ATOM	68	CD1	ILE	9	2.129	65.847	-41.891	1.00	43.63
	ATOM	69	C	ILE	9	5.306	64.530	-38.403	1.00	35.28
	ATOM	70	O	ILE	9	6.522	64.673	-38.495	1.00	33.75
	ATOM	71	N	ALA	10	4.756	63.401	-37.971	1.00	35.78
	ATOM	72	CA	ALA	10	5.596	62.268	-37.617	1.00	35.28
10	ATOM	73	CB	ALA	10	4.755	61.148	-37.050	1.00	28.47
	ATOM	74	C	ALA	10	6.365	61.772	-38.835	1.00	36.49
	ATOM	75	O	ALA	10	5.830	61.704	-39.938	1.00	37.41
	ATOM	76	N	ASP	11	7.628	61.430	-38.628	1.00	39.44
	ATOM	77	CA	ASP	11	8.474	60.931	-39.706	1.00	40.77
15	ATOM	78	CB	ASP	11	9.832	61.633	-39.673	1.00	38.90
	ATOM	79	CG	ASP	11	10.798	61.066	-40.684	1.00	40.34
	ATOM	80	OD1	ASP	11	10.339	60.508	-41.710	1.00	39.50
	ATOM	81	OD2	ASP	11	12.016	61.189	-40.455	1.00	37.38
	ATOM	82	C	ASP	11	8.655	59.423	-39.574	1.00	42.25
20	ATOM	83	O	ASP	11	9.542	58.945	-38.865	1.00	40.54
	ATOM	84	N	SER	12	7.804	58.682	-40.274	1.00	42.92
	ATOM	85	CA	SER	12	7.811	57.221	-40.231	1.00	43.84
	ATOM	86	CB	SER	12	6.628	56.682	-41.028	1.00	42.13
	ATOM	87	OG	SER	12	6.684	57.139	-42.367	1.00	43.31
25	ATOM	88	C	SER	12	9.084	56.565	-40.739	1.00	46.10
	ATOM	89	O	SER	12	9.229	55.345	-40.663	1.00	43.90
	ATOM	90	N	GLU	13	10.008	57.370	-41.249	1.00	46.72
	ATOM	91	CA	GLU	13	11.248	56.829	-41.772	1.00	44.93
	ATOM	92	CB	GLU	13	11.558	57.461	-43.117	1.00	46.71
30	ATOM	93	CG	GLU	13	10.557	57.060	-44.164	1.00	58.25
	ATOM	94	CD	GLU	13	11.125	57.153	-45.565	1.00	66.70
	ATOM	95	OE1	GLU	13	12.160	56.477	-45.822	1.00	74.10
	ATOM	96	OE2	GLU	13	10.548	57.894	-46.406	1.00	71.55
	ATOM	97	C	GLU	13	12.460	56.912	-40.850	1.00	43.24
35	ATOM	98	O	GLU	13	13.594	56.706	-41.289	1.00	44.25
	ATOM	99	N	THR	14	12.233	57.223	-39.583	1.00	37.06
	ATOM	100	CA	THR	14	13.331	57.246	-38.625	1.00	36.77
	ATOM	101	CB	THR	14	13.911	58.673	-38.394	1.00	33.39
	ATOM	102	OG1	THR	14	13.014	59.442	-37.594	1.00	41.69
40	ATOM	103	CG2	THR	14	14.134	59.381	-39.721	1.00	31.66
	ATOM	104	C	THR	14	12.777	56.679	-37.322	1.00	34.96
	ATOM	105	O	THR	14	11.591	56.801	-37.038	1.00	35.04
	ATOM	106	N	PRO	15	13.628	56.031	-36.527	1.00	37.11
	ATOM	107	CD	PRO	15	15.057	55.803	-36.787	1.00	36.50
45	ATOM	108	CA	PRO	15	13.226	55.429	-35.251	1.00	38.81
	ATOM	109	CB	PRO	15	14.497	54.715	-34.790	1.00	39.15
	ATOM	110	CG	PRO	15	15.286	54.518	-36.062	1.00	37.32
	ATOM	111	C	PRO	15	12.770	56.458	-34.225	1.00	37.78
	ATOM	112	O	PRO	15	13.216	57.601	-34.256	1.00	35.91
50	ATOM	113	N	THR	16	11.888	56.060	-33.313	1.00	36.67
	ATOM	114	CA	THR	16	11.440	56.993	-32.292	1.00	38.46
	ATOM	115	CB	THR	16	10.293	56.415	-31.441	1.00	38.61
	ATOM	116	OG1	THR	16	10.758	55.267	-30.723	1.00	42.44
	ATOM	117	CG2	THR	16	9.125	56.017	-32.328	1.00	40.11
55	ATOM	118	C	THR	16	12.637	57.226	-31.395	1.00	35.97
	ATOM	119	O	THR	16	13.354	56.287	-31.076	1.00	39.85
	ATOM	120	N	ILE	17	12.859	58.472	-30.996	1.00	33.66
	ATOM	121	CA	ILE	17	13.992	58.798	-30.142	1.00	34.76
	ATOM	122	CB	ILE	17	14.286	60.303	-30.183	1.00	32.44
60	ATOM	123	CG2	ILE	17	15.387	60.647	-29.199	1.00	28.66
	ATOM	124	CG1	ILE	17	14.666	60.709	-31.612	1.00	33.15
	ATOM	125	CD1	ILE	17	14.904	62.194	-31.799	1.00	31.19
	ATOM	126	C	ILE	17	13.799	58.377	-28.691	1.00	40.64
	ATOM	127	O	ILE	17	12.799	58.709	-28.068	1.00	44.07
65	ATOM	128	N	GLN	18	14.765	57.638	-28.154	1.00	45.91
	ATOM	129	CA	GLN	18	14.704	57.184	-26.766	1.00	50.22

	ATOM	130	CB	GLN	18	15.080	55.710	-26.662	1.00	53.09
	ATOM	131	CG	GLN	18	14.001	54.878	-26.020	1.00	57.60
	ATOM	132	CD	GLN	18	12.788	54.785	-26.909	1.00	61.39
	ATOM	133	OE1	GLN	18	12.827	54.148	-27.965	1.00	68.36
5	ATOM	134	NE2	GLN	18	11.706	55.436	-26.503	1.00	64.93
	ATOM	135	C	GLN	18	15.654	57.988	-25.898	1.00	51.56
	ATOM	136	O	GLN	18	16.815	58.181	-26.247	1.00	56.44
	ATOM	137	N	LYS	19	15.166	58.451	-24.759	1.00	52.60
	ATOM	138	CA	LYS	19	16.011	59.228	-23.868	1.00	56.10
10	ATOM	139	CB	LYS	19	16.328	60.592	-24.482	1.00	55.55
	ATOM	140	CG	LYS	19	17.114	61.467	-23.533	1.00	61.15
	ATOM	141	CD	LYS	19	17.633	62.733	-24.182	1.00	64.42
	ATOM	142	CE	LYS	19	18.435	63.558	-23.164	1.00	64.84
	ATOM	143	NZ	LYS	19	19.009	64.810	-23.753	1.00	69.52
15	ATOM	144	C	LYS	19	15.387	59.427	-22.495	1.00	57.41
	ATOM	145	O	LYS	19	14.205	59.755	-22.385	1.00	61.02
	ATOM	146	N	GLY	20	16.196	59.232	-21.453	1.00	58.84
	ATOM	147	CA	GLY	20	15.726	59.392	-20.090	1.00	56.66
	ATOM	148	C	GLY	20	14.422	58.665	-19.823	1.00	56.87
20	ATOM	149	O	GLY	20	13.572	59.178	-19.094	1.00	58.64
	ATOM	150	N	SER	21	14.270	57.475	-20.407	1.00	54.49
	ATOM	151	CA	SER	21	13.064	56.660	-20.244	1.00	53.68
	ATOM	152	CB	SER	21	12.866	56.285	-18.769	1.00	55.80
	ATOM	153	OG	SER	21	12.187	57.300	-18.054	1.00	65.34
25	ATOM	154	C	SER	21	11.796	57.355	-20.796	1.00	51.41
	ATOM	155	O	SER	21	10.665	57.099	-20.348	1.00	47.74
	ATOM	156	N	TYR	22	12.007	58.238	-21.771	1.00	46.25
	ATOM	157	CA	TYR	22	10.934	58.964	-22.436	1.00	40.57
	ATOM	158	CB	TYR	22	11.092	60.472	-22.229	1.00	43.27
30	ATOM	159	CG	TYR	22	10.447	60.995	-20.972	1.00	49.18
	ATOM	160	CD1	TYR	22	10.826	60.521	-19.721	1.00	52.61
	ATOM	161	CE1	TYR	22	10.220	60.994	-18.553	1.00	51.92
	ATOM	162	CD2	TYR	22	9.445	61.958	-21.030	1.00	50.01
	ATOM	163	CE2	TYR	22	8.836	62.438	-19.870	1.00	50.94
35	ATOM	164	CZ	TYR	22	9.228	61.952	-18.637	1.00	50.78
	ATOM	165	OH	TYR	22	8.624	62.423	-17.492	1.00	53.87
	ATOM	166	C	TYR	22	11.020	58.652	-23.926	1.00	37.85
	ATOM	167	O	TYR	22	12.106	58.409	-24.450	1.00	38.70
	ATOM	168	N	THR	23	9.882	58.637	-24.608	1.00	33.81
40	ATOM	169	CA	THR	23	9.883	58.380	-26.044	1.00	32.40
	ATOM	170	CB	THR	23	8.824	57.330	-26.465	1.00	31.58
	ATOM	171	OG1	THR	23	9.032	56.108	-25.751	1.00	30.76
	ATOM	172	CG2	THR	23	8.924	57.058	-27.951	1.00	24.08
	ATOM	173	C	THR	23	9.552	59.685	-26.747	1.00	33.28
45	ATOM	174	O	THR	23	8.603	60.367	-26.376	1.00	30.44
	ATOM	175	N	PHE	24	10.337	60.029	-27.757	1.00	32.16
	ATOM	176	CA	PHE	24	10.106	61.251	-28.507	1.00	30.68
	ATOM	177	CB	PHE	24	11.311	62.184	-28.398	1.00	28.20
	ATOM	178	CG	PHE	24	11.539	62.711	-27.016	1.00	31.00
50	ATOM	179	CD1	PHE	24	12.247	61.970	-26.085	1.00	31.69
	ATOM	180	CD2	PHE	24	11.016	63.942	-26.633	1.00	31.42
	ATOM	181	CE1	PHE	24	12.429	62.446	-24.792	1.00	32.57
	ATOM	182	CE2	PHE	24	11.194	64.423	-25.347	1.00	25.01
	ATOM	183	CZ	PHE	24	11.902	63.673	-24.426	1.00	30.12
55	ATOM	184	C	PHE	24	9.812	60.964	-29.974	1.00	31.73
	ATOM	185	O	PHE	24	10.536	60.226	-30.635	1.00	33.31
	ATOM	186	N	VAL	25	8.735	61.548	-30.474	1.00	32.92
	ATOM	187	CA	VAL	25	8.353	61.367	-31.858	1.00	30.84
	ATOM	188	CB	VAL	25	6.961	61.962	-32.126	1.00	30.29
60	ATOM	189	CG1	VAL	25	6.599	61.801	-33.581	1.00	28.13
	ATOM	190	CG2	VAL	25	5.938	61.295	-31.248	1.00	29.53
	ATOM	191	C	VAL	25	9.350	62.071	-32.771	1.00	36.08
	ATOM	192	O	VAL	25	9.771	63.192	-32.497	1.00	35.82
	ATOM	193	N	PRO	26	9.762	61.406	-33.860	1.00	38.03
65	ATOM	194	CD	PRO	26	9.569	59.971	-34.117	1.00	38.28
	ATOM	195	CA	PRO	26	10.712	61.978	-34.825	1.00	38.08

	ATOM	196	CB	PRO	26	11.210	60.754	-35.598	1.00	37.62
	ATOM	197	CG	PRO	26	10.889	59.595	-34.709	1.00	36.31
	ATOM	198	C	PRO	26	9.893	62.899	-35.727	1.00	40.85
	ATOM	199	O	PRO	26	9.063	62.428	-36.503	1.00	42.27
5	ATOM	200	N	TRP	27	10.110	64.203	-35.639	1.00	40.86
	ATOM	201	CA	TRP	27	9.314	65.116	-36.451	1.00	37.05
	ATOM	202	CB	TRP	27	9.085	66.432	-35.705	1.00	33.29
	ATOM	203	CG	TRP	27	8.363	66.265	-34.417	1.00	31.06
	ATOM	204	CD2	TRP	27	7.046	65.743	-34.234	1.00	27.79
10	ATOM	205	CE2	TRP	27	6.790	65.739	-32.853	1.00	29.28
	ATOM	206	CE3	TRP	27	6.059	65.275	-35.102	1.00	27.06
	ATOM	207	CD1	TRP	27	8.835	66.553	-33.179	1.00	31.07
	ATOM	208	NE1	TRP	27	7.901	66.241	-32.233	1.00	32.70
	ATOM	209	CZ2	TRP	27	5.584	65.284	-32.313	1.00	25.99
15	ATOM	210	CZ3	TRP	27	4.859	64.823	-34.566	1.00	26.93
	ATOM	211	CH2	TRP	27	4.634	64.831	-33.184	1.00	28.08
	ATOM	212	C	TRP	27	9.861	65.421	-37.826	1.00	38.50
	ATOM	213	O	TRP	27	11.049	65.273	-38.091	1.00	39.07
	ATOM	214	N	LEU	28	8.952	65.847	-38.696	1.00	40.19
20	ATOM	215	CA	LEU	28	9.263	66.227	-40.065	1.00	39.45
	ATOM	216	CB	LEU	28	8.806	65.142	-41.032	1.00	44.18
	ATOM	217	CG	LEU	28	9.398	65.213	-42.438	1.00	46.83
	ATOM	218	CD1	LEU	28	10.899	64.915	-42.357	1.00	45.15
	ATOM	219	CD2	LEU	28	8.692	64.206	-43.350	1.00	49.15
25	ATOM	220	C	LEU	28	8.436	67.486	-40.291	1.00	39.80
	ATOM	221	O	LEU	28	7.294	67.551	-39.860	1.00	42.40
	ATOM	222	N	LEU	29	8.998	68.487	-40.948	1.00	33.58
	ATOM	223	CA	LEU	29	8.243	69.703	-41.173	1.00	29.21
	ATOM	224	CB	LEU	29	9.121	70.782	-41.805	1.00	27.16
30	ATOM	225	CG	LEU	29	8.377	72.080	-42.137	1.00	22.36
	ATOM	226	CD1	LEU	29	8.087	72.843	-40.855	1.00	21.47
	ATOM	227	CD2	LEU	29	9.190	72.922	-43.083	1.00	14.53
	ATOM	228	C	LEU	29	7.027	69.480	-42.065	1.00	29.98
	ATOM	229	O	LEU	29	7.131	68.938	-43.161	1.00	30.36
35	ATOM	230	N	SER	30	5.871	69.903	-41.573	1.00	29.88
	ATOM	231	CA	SER	30	4.638	69.798	-42.330	1.00	28.49
	ATOM	232	CB	SER	30	3.434	69.670	-41.396	1.00	26.13
	ATOM	233	OG	SER	30	2.231	69.664	-42.131	1.00	20.25
	ATOM	234	C	SER	30	4.567	71.102	-43.115	1.00	31.61
40	ATOM	235	O	SER	30	4.416	71.101	-44.332	1.00	35.14
	ATOM	236	N	PHE	31	4.688	72.218	-42.402	1.00	32.12
	ATOM	237	CA	PHE	31	4.673	73.528	-43.029	1.00	30.76
	ATOM	238	CB	PHE	31	3.288	73.841	-43.609	1.00	27.55
	ATOM	239	CG	PHE	31	2.341	74.490	-42.638	1.00	31.23
45	ATOM	240	CD1	PHE	31	2.390	75.860	-42.408	1.00	32.00
	ATOM	241	CD2	PHE	31	1.407	73.728	-41.940	1.00	27.89
	ATOM	242	CE1	PHE	31	1.528	76.459	-41.503	1.00	29.70
	ATOM	243	CE2	PHE	31	0.543	74.322	-41.035	1.00	26.15
	ATOM	244	CZ	PHE	31	0.603	75.688	-40.815	1.00	28.30
50	ATOM	245	C	PHE	31	5.073	74.589	-42.018	1.00	32.06
	ATOM	246	O	PHE	31	4.858	74.439	-40.815	1.00	32.26
	ATOM	247	N	LYS	32	5.673	75.655	-42.516	1.00	34.83
	ATOM	248	CA	LYS	32	6.092	76.754	-41.670	1.00	37.13
	ATOM	249	CB	LYS	32	7.616	76.794	-41.577	1.00	36.55
55	ATOM	250	CG	LYS	32	8.134	77.950	-40.767	1.00	40.62
	ATOM	251	CD	LYS	32	9.624	78.129	-40.924	1.00	44.56
	ATOM	252	CE	LYS	32	10.034	79.480	-40.335	1.00	46.29
	ATOM	253	NZ	LYS	32	11.511	79.695	-40.332	1.00	50.58
	ATOM	254	C	LYS	32	5.560	78.032	-42.310	1.00	36.39
60	ATOM	255	O	LYS	32	5.680	78.224	-43.521	1.00	38.69
	ATOM	256	N	ARG	33	4.953	78.896	-41.504	1.00	37.90
	ATOM	257	CA	ARG	33	4.401	80.143	-42.016	1.00	35.74
	ATOM	258	CB	ARG	33	2.882	80.031	-42.076	1.00	36.79
	ATOM	259	CG	ARG	33	2.153	81.280	-42.516	1.00	31.09
65	ATOM	260	CD	ARG	33	0.692	80.948	-42.794	1.00	31.62
	ATOM	261	NE	ARG	33	-0.056	82.063	-43.370	1.00	38.16

	ATOM	262	CZ	ARG	33	-0.873	82.849	-42.679	1.00	39.15
	ATOM	263	NH1	ARG	33	-1.054	82.645	-41.382	1.00	43.41
	ATOM	264	NH2	ARG	33	-1.508	83.841	-43.280	1.00	40.65
	ATOM	265	C	ARG	33	4.815	81.305	-41.121	1.00	37.71
5	ATOM	266	O	ARG	33	4.495	81.331	-39.929	1.00	41.16
	ATOM	267	N	GLY	34	5.536	82.268	-41.686	1.00	36.59
	ATOM	268	CA	GLY	34	5.962	83.401	-40.892	1.00	36.72
	ATOM	269	C	GLY	34	7.335	83.206	-40.284	1.00	40.01
	ATOM	270	O	GLY	34	8.078	82.301	-40.664	1.00	39.85
10	ATOM	271	N	SER	35	7.659	84.047	-39.311	1.00	38.74
	ATOM	272	CA	SER	35	8.961	84.004	-38.662	1.00	40.23
	ATOM	273	CB	SER	35	9.658	85.340	-38.883	1.00	43.02
	ATOM	274	OG	SER	35	8.817	86.412	-38.458	1.00	48.05
	ATOM	275	C	SER	35	8.952	83.701	-37.162	1.00	40.54
15	ATOM	276	O	SER	35	9.988	83.335	-36.607	1.00	43.74
	ATOM	277	N	ALA	36	7.801	83.849	-36.512	1.00	35.56
	ATOM	278	CA	ALA	36	7.690	83.630	-35.074	1.00	29.62
	ATOM	279	CB	ALA	36	6.287	83.973	-34.619	1.00	25.47
	ATOM	280	C	ALA	36	8.071	82.242	-34.565	1.00	31.23
20	ATOM	281	O	ALA	36	8.451	82.097	-33.410	1.00	29.56
	ATOM	282	N	LEU	37	7.975	81.225	-35.414	1.00	34.09
	ATOM	283	CA	LEU	37	8.289	79.861	-34.999	1.00	32.32
	ATOM	284	CB	LEU	37	7.001	79.051	-34.870	1.00	26.47
	ATOM	285	CG	LEU	37	5.995	79.600	-33.862	1.00	25.72
25	ATOM	286	CD1	LEU	37	4.620	79.042	-34.134	1.00	24.61
	ATOM	287	CD2	LEU	37	6.450	79.269	-32.460	1.00	27.53
	ATOM	288	C	LEU	37	9.240	79.153	-35.948	1.00	35.16
	ATOM	289	O	LEU	37	9.164	79.314	-37.160	1.00	36.47
	ATOM	290	N	GLU	38	10.133	78.358	-35.378	1.00	38.30
30	ATOM	291	CA	GLU	38	11.107	77.609	-36.156	1.00	41.17
	ATOM	292	CB	GLU	38	12.473	78.280	-36.083	1.00	46.15
	ATOM	293	CG	GLU	38	12.739	79.335	-37.131	1.00	50.01
	ATOM	294	CD	GLU	38	13.995	80.137	-36.813	1.00	51.18
	ATOM	295	OE1	GLU	38	14.987	79.519	-36.349	1.00	48.40
35	ATOM	296	OE2	GLU	38	13.984	81.378	-37.028	1.00	50.23
	ATOM	297	C	GLU	38	11.258	76.196	-35.627	1.00	42.69
	ATOM	298	O	GLU	38	10.963	75.917	-34.473	1.00	42.55
	ATOM	299	N	GLU	39	11.725	75.298	-36.478	1.00	44.75
	ATOM	300	CA	GLU	39	11.942	73.927	-36.044	1.00	46.50
40	ATOM	301	CB	GLU	39	11.651	72.951	-37.179	1.00	47.52
	ATOM	302	CG	GLU	39	12.114	73.460	-38.527	1.00	58.50
	ATOM	303	CD	GLU	39	11.949	72.429	-39.626	1.00	62.35
	ATOM	304	OE1	GLU	39	12.112	72.794	-40.820	1.00	64.58
	ATOM	305	OE2	GLU	39	11.666	71.254	-39.294	1.00	63.04
45	ATOM	306	C	GLU	39	13.413	73.888	-35.673	1.00	44.56
	ATOM	307	O	GLU	39	14.244	74.453	-36.373	1.00	48.11
	ATOM	308	N	LYS	40	13.736	73.245	-34.563	1.00	40.71
	ATOM	309	CA	LYS	40	15.116	73.168	-34.125	1.00	38.08
	ATOM	310	CB	LYS	40	15.464	74.384	-33.269	1.00	41.28
50	ATOM	311	CG	LYS	40	16.838	74.308	-32.614	1.00	41.52
	ATOM	312	CD	LYS	40	16.983	75.368	-31.530	1.00	42.73
	ATOM	313	CE	LYS	40	18.277	75.217	-30.750	1.00	41.97
	ATOM	314	NZ	LYS	40	18.332	76.191	-29.616	1.00	45.94
	ATOM	315	C	LYS	40	15.366	71.907	-33.326	1.00	39.66
55	ATOM	316	O	LYS	40	14.834	71.735	-32.232	1.00	36.18
	ATOM	317	N	GLU	41	16.182	71.025	-33.888	1.00	40.16
	ATOM	318	CA	GLU	41	16.541	69.775	-33.234	1.00	42.80
	ATOM	319	CB	GLU	41	17.549	70.069	-32.130	1.00	45.31
	ATOM	320	CG	GLU	41	18.762	70.827	-32.656	1.00	56.07
60	ATOM	321	CD	GLU	41	19.574	71.509	-31.559	1.00	59.42
	ATOM	322	OE1	GLU	41	18.995	72.298	-30.772	1.00	63.71
	ATOM	323	OE2	GLU	41	20.798	71.259	-31.488	1.00	63.71
	ATOM	324	C	GLU	41	15.331	69.040	-32.682	1.00	38.93
	ATOM	325	O	GLU	41	15.284	68.695	-31.504	1.00	39.03
65	ATOM	326	N	ASN	42	14.349	68.824	-33.546	1.00	34.69
	ATOM	327	CA	ASN	42	13.135	68.108	-33.195	1.00	32.96

	ATOM	328	CB	ASN	42	13.484	66.737	-32.628	1.00	32.65
	ATOM	329	CG	ASN	42	12.477	65.696	-33.006	1.00	35.50
	ATOM	330	OD1	ASN	42	12.068	64.888	-32.183	1.00	40.26
	ATOM	331	ND2	ASN	42	12.071	65.702	-34.266	1.00	29.28
5	ATOM	332	C	ASN	42	12.215	68.823	-32.217	1.00	31.44
	ATOM	333	O	ASN	42	11.345	68.204	-31.612	1.00	22.78
	ATOM	334	N	LYS	43	12.402	70.125	-32.073	1.00	33.26
	ATOM	335	CA	LYS	43	11.583	70.914	-31.169	1.00	33.68
	ATOM	336	CB	LYS	43	12.378	71.277	-29.915	1.00	33.55
10	ATOM	337	CG	LYS	43	12.710	70.094	-29.041	1.00	40.49
	ATOM	338	CD	LYS	43	13.573	70.508	-27.877	1.00	45.39
	ATOM	339	CE	LYS	43	14.928	70.990	-28.357	1.00	52.57
	ATOM	340	NZ	LYS	43	15.825	71.300	-27.214	1.00	62.06
	ATOM	341	C	LYS	43	11.138	72.181	-31.863	1.00	32.05
15	ATOM	342	O	LYS	43	11.665	72.534	-32.909	1.00	37.32
	ATOM	343	N	ILE	44	10.155	72.862	-31.293	1.00	31.31
	ATOM	344	CA	ILE	44	9.699	74.100	-31.892	1.00	29.52
	ATOM	345	CB	ILE	44	8.173	74.207	-31.864	1.00	26.38
	ATOM	346	CG2	ILE	44	7.743	75.533	-32.455	1.00	26.09
20	ATOM	347	CG1	ILE	44	7.568	73.064	-32.680	1.00	28.61
	ATOM	348	CD1	ILE	44	6.072	73.075	-32.749	1.00	21.51
	ATOM	349	C	ILE	44	10.316	75.262	-31.134	1.00	31.60
	ATOM	350	O	ILE	44	10.152	75.387	-29.922	1.00	34.50
	ATOM	351	N	LEU	45	11.049	76.098	-31.856	1.00	31.44
25	ATOM	352	CA	LEU	45	11.707	77.255	-31.268	1.00	31.40
	ATOM	353	CB	LEU	45	13.100	77.414	-31.872	1.00	33.60
	ATOM	354	CG	LEU	45	13.879	78.655	-31.443	1.00	34.36
	ATOM	355	CD1	LEU	45	14.225	78.573	-29.970	1.00	29.00
	ATOM	356	CD2	LEU	45	15.136	78.760	-32.285	1.00	32.42
30	ATOM	357	C	LEU	45	10.911	78.541	-31.476	1.00	27.80
	ATOM	358	O	LEU	45	10.508	78.864	-32.588	1.00	27.94
	ATOM	359	N	VAL	46	10.692	79.269	-30.388	1.00	28.00
	ATOM	360	CA	VAL	46	9.961	80.527	-30.432	1.00	31.53
	ATOM	361	CB	VAL	46	9.275	80.785	-29.080	1.00	27.89
35	ATOM	362	CG1	VAL	46	8.472	82.062	-29.131	1.00	26.10
	ATOM	363	CG2	VAL	46	8.391	79.619	-28.733	1.00	23.39
	ATOM	364	C	VAL	46	10.934	81.675	-30.742	1.00	33.94
	ATOM	365	O	VAL	46	11.926	81.861	-30.035	1.00	39.68
	ATOM	366	N	LYS	47	10.648	82.442	-31.791	1.00	33.59
40	ATOM	367	CA	LYS	47	11.516	83.553	-32.174	1.00	37.05
	ATOM	368	CB	LYS	47	11.836	83.483	-33.671	1.00	34.49
	ATOM	369	CG	LYS	47	12.292	82.111	-34.135	1.00	37.53
	ATOM	370	CD	LYS	47	13.516	81.635	-33.364	1.00	41.69
	ATOM	371	CE	LYS	47	14.799	81.964	-34.105	1.00	42.95
45	ATOM	372	NZ	LYS	47	14.803	83.363	-34.590	1.00	44.50
	ATOM	373	C	LYS	47	10.933	84.926	-31.838	1.00	37.34
	ATOM	374	O	LYS	47	11.630	85.934	-31.903	1.00	43.04
	ATOM	375	N	GLU	48	9.656	84.958	-31.483	1.00	36.99
	ATOM	376	CA	GLU	48	8.976	86.197	-31.115	1.00	37.31
50	ATOM	377	CB	GLU	48	8.057	86.668	-32.229	1.00	36.53
	ATOM	378	CG	GLU	48	8.746	87.155	-33.469	1.00	46.74
	ATOM	379	CD	GLU	48	7.745	87.557	-34.537	1.00	52.39
	ATOM	380	OE1	GLU	48	6.707	88.177	-34.174	1.00	51.33
	ATOM	381	OE2	GLU	48	7.996	87.259	-35.733	1.00	54.55
55	ATOM	382	C	GLU	48	8.120	85.888	-29.911	1.00	36.94
	ATOM	383	O	GLU	48	7.346	84.941	-29.947	1.00	40.21
	ATOM	384	N	THR	49	8.225	86.674	-28.848	1.00	36.34
	ATOM	385	CA	THR	49	7.413	86.365	-27.682	1.00	35.48
	ATOM	386	CB	THR	49	7.982	87.010	-26.388	1.00	31.16
60	ATOM	387	OG1	THR	49	7.295	88.230	-26.117	1.00	32.48
	ATOM	388	CG2	THR	49	9.457	87.295	-26.534	1.00	30.36
	ATOM	389	C	THR	49	5.963	86.803	-27.905	1.00	34.59
	ATOM	390	O	THR	49	5.691	87.745	-28.647	1.00	35.13
	ATOM	391	N	GLY	50	5.038	86.088	-27.271	1.00	32.49
65	ATOM	392	CA	GLY	50	3.632	86.405	-27.406	1.00	28.17
	ATOM	393	C	GLY	50	2.772	85.253	-26.941	1.00	29.99

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	ATOM	394	O	GLY	50	3.239	84.382	-26.225	1.00	29.76
	ATOM	395	N	TYR	51	1.506	85.255	-27.345	1.00	31.68
	ATOM	396	CA	TYR	51	0.575	84.192	-26.984	1.00	32.88
	ATOM	397	CB	TYR	51	-0.784	84.776	-26.582	1.00	35.97
5	ATOM	398	CG	TYR	51	-0.743	85.497	-25.267	1.00	43.03
	ATOM	399	CD1	TYR	51	-0.278	86.812	-25.182	1.00	44.15
	ATOM	400	CE1	TYR	51	-0.151	87.455	-23.950	1.00	47.81
	ATOM	401	CD2	TYR	51	-1.089	84.838	-24.089	1.00	47.08
	ATOM	402	CE2	TYR	51	-0.963	85.465	-22.850	1.00	51.79
10	ATOM	403	CZ	TYR	51	-0.490	86.773	-22.784	1.00	51.15
	ATOM	404	OH	TYR	51	-0.318	87.374	-21.551	1.00	52.20
	ATOM	405	C	TYR	51	0.394	83.217	-28.140	1.00	31.97
	ATOM	406	O	TYR	51	0.077	83.611	-29.259	1.00	30.01
	ATOM	407	N	PHE	52	0.600	81.934	-27.862	1.00	31.50
15	ATOM	408	CA	PHE	52	0.470	80.919	-28.893	1.00	30.41
	ATOM	409	CB	PHE	52	1.815	80.233	-29.143	1.00	31.30
	ATOM	410	CG	PHE	52	2.908	81.159	-29.576	1.00	29.70
	ATOM	411	CD1	PHE	52	3.501	82.027	-28.674	1.00	26.20
	ATOM	412	CD2	PHE	52	3.352	81.154	-30.890	1.00	28.38
20	ATOM	413	CE1	PHE	52	4.515	82.872	-29.075	1.00	27.17
	ATOM	414	CE2	PHE	52	4.368	81.998	-31.299	1.00	27.76
	ATOM	415	CZ	PHE	52	4.950	82.857	-30.392	1.00	27.27
	ATOM	416	C	PHE	52	-0.542	79.842	-28.560	1.00	27.65
	ATOM	417	O	PHE	52	-0.732	79.486	-27.404	1.00	24.85
25	ATOM	418	N	PHE	53	-1.193	79.337	-29.597	1.00	27.18
	ATOM	419	CA	PHE	53	-2.142	78.244	-29.464	1.00	27.05
	ATOM	420	CB	PHE	53	-3.279	78.383	-30.468	1.00	28.76
	ATOM	421	CG	PHE	53	-4.186	77.196	-30.517	1.00	28.00
	ATOM	422	CD1	PHE	53	-5.049	76.918	-29.469	1.00	26.49
30	ATOM	423	CD2	PHE	53	-4.166	76.340	-31.609	1.00	28.04
	ATOM	424	CE1	PHE	53	-5.880	75.804	-29.508	1.00	26.54
	ATOM	425	CE2	PHE	53	-4.995	75.223	-31.651	1.00	25.78
	ATOM	426	CZ	PHE	53	-5.851	74.958	-30.598	1.00	24.39
	ATOM	427	C	PHE	53	-1.288	77.032	-29.825	1.00	27.56
35	ATOM	428	O	PHE	53	-0.738	76.968	-30.920	1.00	25.39
	ATOM	429	N	ILE	54	-1.165	76.086	-28.900	1.00	28.87
	ATOM	430	CA	ILE	54	-0.342	74.901	-29.118	1.00	28.95
	ATOM	431	CB	ILE	54	0.736	74.807	-28.018	1.00	29.60
	ATOM	432	CG2	ILE	54	1.764	73.749	-28.387	1.00	27.42
40	ATOM	433	CG1	ILE	54	1.420	76.167	-27.857	1.00	26.41
	ATOM	434	CD1	ILE	54	2.246	76.306	-26.616	1.00	28.70
	ATOM	435	C	ILE	54	-1.185	73.627	-29.127	1.00	28.57
	ATOM	436	O	ILE	54	-2.075	73.453	-28.301	1.00	29.17
	ATOM	437	N	TYR	55	-0.904	72.727	-30.061	1.00	26.97
45	ATOM	438	CA	TYR	55	-1.683	71.499	-30.157	1.00	24.22
	ATOM	439	CB	TYR	55	-2.787	71.674	-31.201	1.00	21.82
	ATOM	440	CG	TYR	55	-2.286	72.019	-32.582	1.00	27.68
	ATOM	441	CD1	TYR	55	-1.989	71.023	-33.507	1.00	27.01
	ATOM	442	CE1	TYR	55	-1.509	71.341	-34.765	1.00	27.44
50	ATOM	443	CD2	TYR	55	-2.089	73.344	-32.957	1.00	27.16
	ATOM	444	CE2	TYR	55	-1.606	73.669	-34.215	1.00	25.39
	ATOM	445	CZ	TYR	55	-1.318	72.666	-35.111	1.00	29.02
	ATOM	446	OH	TYR	55	-0.829	72.990	-36.351	1.00	31.82
	ATOM	447	C	TYR	55	-0.844	70.284	-30.488	1.00	25.16
55	ATOM	448	O	TYR	55	0.275	70.406	-30.934	1.00	28.47
	ATOM	449	N	GLY	56	-1.388	69.102	-30.254	1.00	27.44
	ATOM	450	CA	GLY	56	-0.650	67.892	-30.543	1.00	22.07
	ATOM	451	C	GLY	56	-1.519	66.657	-30.459	1.00	25.83
	ATOM	452	O	GLY	56	-2.444	66.596	-29.655	1.00	26.00
60	ATOM	453	N	GLN	57	-1.233	65.680	-31.310	1.00	22.56
	ATOM	454	CA	GLN	57	-1.965	64.424	-31.326	1.00	20.91
	ATOM	455	CB	GLN	57	-3.063	64.420	-32.391	1.00	17.87
	ATOM	456	CG	GLN	57	-3.754	63.067	-32.515	1.00	16.86
	ATOM	457	CD	GLN	57	-4.901	63.056	-33.503	1.00	22.06
65	ATOM	458	OE1	GLN	57	-5.905	63.733	-33.310	1.00	27.26
	ATOM	459	NE2	GLN	57	-4.758	62.277	-34.568	1.00	20.20

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	ATOM	460	C	GLN	57	-0.998	63.294	-31.609	1.00	24.23
	ATOM	461	O	GLN	57	-0.048	63.462	-32.371	1.00	25.16
	ATOM	462	N	VAL	58	-1.251	62.148	-30.984	1.00	24.62
	ATOM	463	CA	VAL	58	-0.429	60.954	-31.145	1.00	26.68
5	ATOM	464	CB	VAL	58	0.534	60.765	-29.947	1.00	25.55
	ATOM	465	CG1	VAL	58	1.218	59.423	-30.035	1.00	15.69
	ATOM	466	CG2	VAL	58	1.566	61.872	-29.918	1.00	25.77
	ATOM	467	C	VAL	58	-1.357	59.758	-31.185	1.00	31.06
	ATOM	468	O	VAL	58	-2.330	59.724	-30.443	1.00	27.74
10	ATOM	469	N	LEU	59	-1.066	58.784	-32.048	1.00	33.82
	ATOM	470	CA	LEU	59	-1.873	57.562	-32.133	1.00	29.85
	ATOM	471	CB	LEU	59	-2.045	57.105	-33.579	1.00	29.62
	ATOM	472	CG	LEU	59	-3.254	56.216	-33.898	1.00	28.11
	ATOM	473	CD1	LEU	59	-3.062	55.588	-35.264	1.00	26.67
15	ATOM	474	CD2	LEU	59	-3.437	55.142	-32.867	1.00	22.45
	ATOM	475	C	LEU	59	-1.145	56.468	-31.357	1.00	30.98
	ATOM	476	O	LEU	59	-0.054	56.047	-31.737	1.00	32.52
	ATOM	477	N	TYR	60	-1.750	56.012	-30.265	1.00	33.01
	ATOM	478	CA	TYR	60	-1.143	54.970	-29.448	1.00	34.51
20	ATOM	479	CB	TYR	60	-1.470	55.203	-27.977	1.00	40.10
	ATOM	480	CG	TYR	60	-0.923	56.500	-27.467	1.00	45.37
	ATOM	481	CD1	TYR	60	-1.761	57.590	-27.227	1.00	47.13
	ATOM	482	CE1	TYR	60	-1.239	58.816	-26.837	1.00	50.28
	ATOM	483	CD2	TYR	60	0.445	56.667	-27.297	1.00	44.35
25	ATOM	484	CE2	TYR	60	0.979	57.880	-26.910	1.00	48.81
	ATOM	485	CZ	TYR	60	0.137	58.950	-26.687	1.00	51.91
	ATOM	486	OH	TYR	60	0.683	60.162	-26.345	1.00	53.77
	ATOM	487	C	TYR	60	-1.573	53.569	-29.847	1.00	35.49
	ATOM	488	O	TYR	60	-2.756	53.245	-29.862	1.00	36.13
30	ATOM	489	N	THR	61	-0.591	52.741	-30.173	1.00	36.19
	ATOM	490	CA	THR	61	-0.843	51.365	-30.557	1.00	35.36
	ATOM	491	CB	THR	61	-0.364	51.102	-31.986	1.00	34.22
	ATOM	492	OG1	THR	61	1.001	51.511	-32.114	1.00	34.22
	ATOM	493	CG2	THR	61	-1.217	51.867	-32.980	1.00	25.45
35	ATOM	494	C	THR	61	-0.072	50.489	-29.579	1.00	37.82
	ATOM	495	O	THR	61	0.414	49.416	-29.921	1.00	43.70
	ATOM	496	N	ASP	62	0.029	50.981	-28.351	1.00	37.64
	ATOM	497	CA	ASP	62	0.727	50.317	-27.266	1.00	38.08
	ATOM	498	CB	ASP	62	1.647	51.340	-26.607	1.00	38.42
40	ATOM	499	CG	ASP	62	2.564	50.734	-25.583	1.00	41.25
	ATOM	500	OD1	ASP	62	3.741	51.162	-25.525	1.00	38.28
	ATOM	501	OD2	ASP	62	2.102	49.846	-24.838	1.00	41.37
	ATOM	502	C	ASP	62	-0.331	49.822	-26.284	1.00	40.90
	ATOM	503	O	ASP	62	-1.314	50.513	-26.053	1.00	46.73
45	ATOM	504	N	LYS	63	-0.152	48.639	-25.703	1.00	42.62
	ATOM	505	CA	LYS	63	-1.163	48.130	-24.775	1.00	40.92
	ATOM	506	CB	LYS	63	-1.422	46.645	-25.033	1.00	42.79
	ATOM	507	CG	LYS	63	-0.208	45.752	-24.896	1.00	45.91
	ATOM	508	CD	LYS	63	-0.530	44.339	-25.384	1.00	46.22
50	ATOM	509	CE	LYS	63	-0.914	44.343	-26.876	1.00	49.97
	ATOM	510	NZ	LYS	63	-1.458	43.031	-27.371	1.00	49.98
	ATOM	511	C	LYS	63	-0.864	48.347	-23.295	1.00	42.59
	ATOM	512	O	LYS	63	-1.443	47.686	-22.442	1.00	42.46
	ATOM	513	N	THR	64	0.007	49.303	-22.999	1.00	44.06
55	ATOM	514	CA	THR	64	0.393	49.611	-21.629	1.00	44.94
	ATOM	515	CB	THR	64	1.628	50.547	-21.620	1.00	43.79
	ATOM	516	OG1	THR	64	2.680	49.947	-22.382	1.00	46.20
	ATOM	517	CG2	THR	64	2.134	50.777	-20.200	1.00	42.80
	ATOM	518	C	THR	64	-0.698	50.219	-20.732	1.00	45.82
60	ATOM	519	O	THR	64	-0.440	51.160	-20.004	1.00	53.10
	ATOM	520	N	TYR	65	-1.906	49.679	-20.773	1.00	45.11
	ATOM	521	CA	TYR	65	-3.022	50.144	-19.935	1.00	45.72
	ATOM	522	CB	TYR	65	-2.838	49.645	-18.489	1.00	40.46
	ATOM	523	CG	TYR	65	-2.156	50.606	-17.540	1.00	41.90
65	ATOM	524	CD1	TYR	65	-2.886	51.564	-16.835	1.00	42.47
	ATOM	525	CE1	TYR	65	-2.255	52.465	-15.959	1.00	41.83

	ATOM	526	CD2	TYR	65	-0.770	50.566	-17.350	1.00	41.65
	ATOM	527	CE2	TYR	65	-0.126	51.463	-16.480	1.00	40.28
	ATOM	528	CZ	TYR	65	-0.877	52.406	-15.791	1.00	41.90
	ATOM	529	OH	TYR	65	-0.253	53.284	-14.940	1.00	43.84
5	ATOM	530	C	TYR	65	-3.336	51.649	-19.924	1.00	45.31
	ATOM	531	O	TYR	65	-4.495	52.046	-19.746	1.00	49.74
	ATOM	532	N	ALA	66	-2.321	52.483	-20.101	1.00	42.32
	ATOM	533	CA	ALA	66	-2.511	53.928	-20.107	1.00	39.57
	ATOM	534	CB	ALA	66	-2.587	54.452	-18.690	1.00	35.36
10	ATOM	535	C	ALA	66	-1.366	54.593	-20.849	1.00	39.85
	ATOM	536	O	ALA	66	-0.199	54.430	-20.489	1.00	37.60
	ATOM	537	N	MET	67	-1.703	55.328	-21.900	1.00	41.88
	ATOM	538	CA	MET	67	-0.702	56.025	-22.694	1.00	41.48
	ATOM	539	CB	MET	67	-0.637	55.454	-24.116	1.00	39.98
15	ATOM	540	CG	MET	67	-0.081	54.039	-24.214	1.00	37.31
	ATOM	541	SD	MET	67	1.590	53.891	-23.568	1.00	35.46
	ATOM	542	CE	MET	67	2.568	54.495	-24.945	1.00	34.06
	ATOM	543	C	MET	67	-1.047	57.503	-22.747	1.00	39.50
	ATOM	544	O	MET	67	-2.157	57.907	-22.393	1.00	38.44
20	ATOM	545	N	GLY	68	-0.088	58.307	-23.188	1.00	39.84
	ATOM	546	CA	GLY	68	-0.312	59.735	-23.281	1.00	37.61
	ATOM	547	C	GLY	68	0.961	60.492	-23.573	1.00	33.76
	ATOM	548	O	GLY	68	2.042	59.915	-23.575	1.00	35.58
	ATOM	549	N	HIS	69	0.834	61.786	-23.835	1.00	35.07
25	ATOM	550	CA	HIS	69	1.995	62.613	-24.119	1.00	33.70
	ATOM	551	CB	HIS	69	2.142	62.855	-25.633	1.00	37.02
	ATOM	552	CG	HIS	69	0.917	63.404	-26.296	1.00	34.65
	ATOM	553	CD2	HIS	69	0.635	64.644	-26.756	1.00	35.63
	ATOM	554	ND1	HIS	69	-0.194	62.636	-26.569	1.00	37.68
30	ATOM	555	CE1	HIS	69	-1.107	63.378	-27.168	1.00	32.64
	ATOM	556	NE2	HIS	69	-0.628	64.601	-27.295	1.00	35.90
	ATOM	557	C	HIS	69	1.929	63.943	-23.382	1.00	33.98
	ATOM	558	O	HIS	69	0.909	64.292	-22.810	1.00	30.94
	ATOM	559	N	LEU	70	3.032	64.678	-23.396	1.00	35.35
35	ATOM	560	CA	LEU	70	3.104	65.970	-22.730	1.00	30.76
	ATOM	561	CB	LEU	70	4.097	65.915	-21.574	1.00	29.59
	ATOM	562	CG	LEU	70	4.142	64.671	-20.693	1.00	30.71
	ATOM	563	CD1	LEU	70	5.357	64.741	-19.800	1.00	25.60
	ATOM	564	CD2	LEU	70	2.872	64.564	-19.882	1.00	27.39
40	ATOM	565	C	LEU	70	3.599	67.036	-23.703	1.00	33.19
	ATOM	566	O	LEU	70	4.513	66.790	-24.490	1.00	37.07
	ATOM	567	N	ILE	71	2.984	68.212	-23.673	1.00	30.93
	ATOM	568	CA	ILE	71	3.456	69.303	-24.509	1.00	27.94
	ATOM	569	CB	ILE	71	2.308	70.120	-25.108	1.00	28.22
45	ATOM	570	CG2	ILE	71	2.834	71.423	-25.674	1.00	24.54
	ATOM	571	CG1	ILE	71	1.639	69.318	-26.218	1.00	26.34
	ATOM	572	CD1	ILE	71	0.377	69.946	-26.739	1.00	37.50
	ATOM	573	C	ILE	71	4.213	70.132	-23.483	1.00	30.32
	ATOM	574	O	ILE	71	3.615	70.646	-22.546	1.00	23.07
50	ATOM	575	N	GLN	72	5.530	70.233	-23.640	1.00	29.74
	ATOM	576	CA	GLN	72	6.327	70.967	-22.668	1.00	27.85
	ATOM	577	CB	GLN	72	7.385	70.040	-22.090	1.00	26.70
	ATOM	578	CG	GLN	72	6.810	68.737	-21.613	1.00	28.28
	ATOM	579	CD	GLN	72	7.814	67.908	-20.865	1.00	33.11
55	ATOM	580	OE1	GLN	72	8.900	67.635	-21.369	1.00	41.22
	ATOM	581	NE2	GLN	72	7.460	67.495	-19.656	1.00	30.18
	ATOM	582	C	GLN	72	6.988	72.231	-23.176	1.00	29.85
	ATOM	583	O	GLN	72	7.205	72.401	-24.377	1.00	28.20
	ATOM	584	N	ARG	73	7.319	73.106	-22.234	1.00	30.90
60	ATOM	585	CA	ARG	73	7.967	74.371	-22.540	1.00	30.79
	ATOM	586	CB	ARG	73	7.042	75.518	-22.160	1.00	29.18
	ATOM	587	CG	ARG	73	7.645	76.877	-22.351	1.00	28.66
	ATOM	588	CD	ARG	73	6.955	77.877	-21.472	1.00	29.25
	ATOM	589	NE	ARG	73	7.482	79.218	-21.669	1.00	35.78
65	ATOM	590	CZ	ARG	73	7.321	80.214	-20.810	1.00	35.27
	ATOM	591	NH1	ARG	73	6.650	80.018	-19.684	1.00	34.95

	ATOM	592	NH2	ARG	73	7.828	81.402	-21.083	1.00	33.33
	ATOM	593	C	ARG	73	9.289	74.528	-21.793	1.00	32.42
	ATOM	594	O	ARG	73	9.340	74.356	-20.580	1.00	34.19
	ATOM	595	N	LYS	74	10.355	74.835	-22.525	1.00	35.90
5	ATOM	596	CA	LYS	74	11.667	75.065	-21.919	1.00	38.57
	ATOM	597	CB	LYS	74	12.787	74.442	-22.758	1.00	43.52
	ATOM	598	CG	LYS	74	12.865	72.926	-22.693	1.00	52.89
	ATOM	599	CD	LYS	74	13.985	72.359	-23.596	1.00	58.14
	ATOM	600	CE	LYS	74	14.028	70.832	-23.521	1.00	56.75
10	ATOM	601	NZ	LYS	74	15.071	70.251	-24.417	1.00	63.98
	ATOM	602	C	LYS	74	11.880	76.573	-21.873	1.00	40.11
	ATOM	603	O	LYS	74	12.133	77.201	-22.909	1.00	37.55
	ATOM	604	N	LYS	75	11.777	77.150	-20.680	1.00	34.48
	ATOM	605	CA	LYS	75	11.948	78.588	-20.508	1.00	34.24
15	ATOM	606	CB	LYS	75	11.572	78.990	-19.080	1.00	36.16
	ATOM	607	CG	LYS	75	10.162	78.630	-18.618	1.00	35.99
	ATOM	608	CD	LYS	75	9.965	79.091	-17.167	1.00	40.91
	ATOM	609	CE	LYS	75	8.565	78.785	-16.652	1.00	46.16
	ATOM	610	NZ	LYS	75	8.330	79.163	-15.215	1.00	46.03
20	ATOM	611	C	LYS	75	13.394	79.019	-20.775	1.00	30.92
	ATOM	612	O	LYS	75	14.320	78.334	-20.383	1.00	27.47
	ATOM	613	N	VAL	76	13.575	80.159	-21.436	1.00	32.82
	ATOM	614	CA	VAL	76	14.913	80.690	-21.716	1.00	31.51
	ATOM	615	CB	VAL	76	14.912	81.789	-22.794	1.00	29.19
25	ATOM	616	CG1	VAL	76	16.170	81.700	-23.610	1.00	27.09
	ATOM	617	CG2	VAL	76	13.700	81.689	-23.650	1.00	34.94
	ATOM	618	C	VAL	76	15.406	81.370	-20.455	1.00	33.40
	ATOM	619	O	VAL	76	16.574	81.271	-20.095	1.00	32.92
	ATOM	620	N	HIS	77	14.493	82.085	-19.804	1.00	34.88
30	ATOM	621	CA	HIS	77	14.789	82.809	-18.582	1.00	34.59
	ATOM	622	CB	HIS	77	14.241	84.237	-18.671	1.00	35.92
	ATOM	623	CG	HIS	77	14.777	85.019	-19.830	1.00	34.17
	ATOM	624	CD2	HIS	77	14.262	86.069	-20.509	1.00	31.44
	ATOM	625	ND1	HIS	77	16.008	84.764	-20.399	1.00	35.05
35	ATOM	626	CE1	HIS	77	16.229	85.619	-21.379	1.00	29.84
	ATOM	627	NE2	HIS	77	15.186	86.423	-21.465	1.00	38.35
	ATOM	628	C	HIS	77	14.171	82.073	-17.404	1.00	35.63
	ATOM	629	O	HIS	77	13.043	81.587	-17.497	1.00	40.24
	ATOM	630	N	VAL	78	14.896	82.013	-16.289	1.00	37.37
40	ATOM	631	CA	VAL	78	14.404	81.281	-15.136	1.00	41.32
	ATOM	632	CB	VAL	78	15.339	80.093	-14.837	1.00	43.10
	ATOM	633	CG1	VAL	78	14.761	79.243	-13.732	1.00	45.01
	ATOM	634	CG2	VAL	78	15.499	79.239	-16.082	1.00	44.98
	ATOM	635	C	VAL	78	14.075	82.018	-13.831	1.00	43.02
45	ATOM	636	O	VAL	78	12.907	82.069	-13.450	1.00	49.79
	ATOM	637	N	PHE	79	15.053	82.573	-13.127	1.00	41.33
	ATOM	638	CA	PHE	79	14.745	83.270	-11.854	1.00	45.00
	ATOM	639	CB	PHE	79	13.474	84.138	-11.936	1.00	40.51
	ATOM	640	CG	PHE	79	13.480	85.143	-13.037	1.00	39.52
50	ATOM	641	CD1	PHE	79	12.714	84.939	-14.179	1.00	36.80
	ATOM	642	CD2	PHE	79	14.245	86.298	-12.934	1.00	37.89
	ATOM	643	CE1	PHE	79	12.708	85.870	-15.204	1.00	37.52
	ATOM	644	CE2	PHE	79	14.244	87.235	-13.955	1.00	35.59
	ATOM	645	CZ	PHE	79	13.475	87.022	-15.093	1.00	37.58
55	ATOM	646	C	PHE	79	14.522	82.342	-10.660	1.00	43.50
	ATOM	647	O	PHE	79	13.660	81.462	-10.690	1.00	37.85
	ATOM	648	N	GLY	80	15.283	82.579	-9.597	1.00	46.65
	ATOM	649	CA	GLY	80	15.155	81.806	-8.378	1.00	48.38
	ATOM	650	C	GLY	80	15.074	80.301	-8.509	1.00	47.89
60	ATOM	651	O	GLY	80	15.881	79.680	-9.207	1.00	51.15
	ATOM	652	N	ASP	81	14.091	79.713	-7.831	1.00	45.07
	ATOM	653	CA	ASP	81	13.916	78.272	-7.839	1.00	43.99
	ATOM	654	CB	ASP	81	13.541	77.781	-6.434	1.00	45.66
	ATOM	655	CG	ASP	81	12.146	78.214	-6.009	1.00	47.49
65	ATOM	656	OD1	ASP	81	11.554	79.104	-6.659	1.00	51.90
	ATOM	657	OD2	ASP	81	11.640	77.667	-5.009	1.00	48.08

	ATOM	658	C	ASP	81	12.903	77.772	-8.853	1.00	42.29
	ATOM	659	O	ASP	81	12.348	76.685	-8.691	1.00	39.94
	ATOM	660	N	GLU	82	12.658	78.560	-9.895	1.00	39.91
	ATOM	661	CA	GLU	82	11.725	78.150	-10.943	1.00	36.58
5	ATOM	662	CB	GLU	82	11.556	79.235	-12.008	1.00	34.91
	ATOM	663	CG	GLU	82	10.654	80.382	-11.686	1.00	37.82
	ATOM	664	CD	GLU	82	10.107	81.026	-12.950	1.00	40.75
	ATOM	665	OE1	GLU	82	10.882	81.223	-13.907	1.00	38.15
	ATOM	666	OE2	GLU	82	8.898	81.334	-12.995	1.00	47.99
10	ATOM	667	C	GLU	82	12.285	76.931	-11.663	1.00	36.58
	ATOM	668	O	GLU	82	13.497	76.773	-11.779	1.00	38.40
	ATOM	669	N	LEU	83	11.398	76.072	-12.146	1.00	33.83
	ATOM	670	CA	LEU	83	11.809	74.914	-12.923	1.00	32.70
	ATOM	671	CB	LEU	83	10.781	73.789	-12.808	1.00	30.78
15	ATOM	672	CG	LEU	83	10.994	72.732	-11.727	1.00	29.26
	ATOM	673	CD1	LEU	83	11.420	73.368	-10.428	1.00	29.60
	ATOM	674	CD2	LEU	83	9.716	71.953	-11.547	1.00	28.65
	ATOM	675	C	LEU	83	11.801	75.475	-14.329	1.00	33.88
	ATOM	676	O	LEU	83	10.849	76.144	-14.714	1.00	38.46
20	ATOM	677	N	SER	84	12.854	75.228	-15.095	1.00	34.94
	ATOM	678	CA	SER	84	12.911	75.763	-16.446	1.00	39.77
	ATOM	679	CB	SER	84	14.360	75.878	-16.914	1.00	38.92
	ATOM	680	OG	SER	84	14.991	74.621	-16.879	1.00	43.11
	ATOM	681	C	SER	84	12.102	74.949	-17.452	1.00	39.89
25	ATOM	682	O	SER	84	11.879	75.380	-18.583	1.00	42.52
	ATOM	683	N	LEU	85	11.658	73.774	-17.034	1.00	40.92
	ATOM	684	CA	LEU	85	10.865	72.915	-17.893	1.00	34.66
	ATOM	685	CB	LEU	85	11.475	71.514	-17.936	1.00	35.12
	ATOM	686	CG	LEU	85	11.018	70.470	-18.965	1.00	36.92
30	ATOM	687	CD1	LEU	85	9.602	70.034	-18.675	1.00	38.24
	ATOM	688	CD2	LEU	85	11.135	71.045	-20.368	1.00	36.08
	ATOM	689	C	LEU	85	9.465	72.870	-17.310	1.00	35.30
	ATOM	690	O	LEU	85	9.253	72.389	-16.200	1.00	32.85
	ATOM	691	N	VAL	86	8.510	73.402	-18.057	1.00	33.26
35	ATOM	692	CA	VAL	86	7.124	73.420	-17.619	1.00	34.57
	ATOM	693	CB	VAL	86	6.528	74.858	-17.641	1.00	35.39
	ATOM	694	CG1	VAL	86	5.041	74.816	-17.346	1.00	31.44
	ATOM	695	CG2	VAL	86	7.232	75.729	-16.620	1.00	38.08
	ATOM	696	C	VAL	86	6.300	72.567	-18.554	1.00	32.85
40	ATOM	697	O	VAL	86	6.429	72.671	-19.771	1.00	35.10
	ATOM	698	N	THR	87	5.459	71.711	-18.001	1.00	33.41
	ATOM	699	CA	THR	87	4.617	70.916	-18.861	1.00	33.56
	ATOM	700	CB	THR	87	4.623	69.412	-18.445	1.00	32.70
	ATOM	701	OG1	THR	87	3.303	68.864	-18.550	1.00	34.29
45	ATOM	702	CG2	THR	87	5.168	69.243	-17.052	1.00	34.95
	ATOM	703	C	THR	87	3.199	71.501	-18.896	1.00	30.57
	ATOM	704	O	THR	87	2.508	71.568	-17.883	1.00	28.08
	ATOM	705	N	LEU	88	2.820	71.985	-20.075	1.00	28.42
	ATOM	706	CA	LEU	88	1.501	72.540	-20.315	1.00	31.31
50	ATOM	707	CB	LEU	88	1.545	73.620	-21.405	1.00	28.21
	ATOM	708	CG	LEU	88	2.431	74.866	-21.509	1.00	29.12
	ATOM	709	CD1	LEU	88	3.212	75.087	-20.262	1.00	31.12
	ATOM	710	CD2	LEU	88	3.348	74.719	-22.699	1.00	28.72
	ATOM	711	C	LEU	88	0.657	71.378	-20.868	1.00	41.15
55	ATOM	712	O	LEU	88	1.171	70.483	-21.544	1.00	52.30
	ATOM	713	N	PHE	89	-0.630	71.356	-20.578	1.00	39.23
	ATOM	714	CA	PHE	89	-1.475	70.313	-21.164	1.00	42.27
	ATOM	715	CB	PHE	89	-1.474	70.523	-22.681	1.00	35.63
	ATOM	716	CG	PHE	89	-1.566	71.977	-23.060	1.00	35.06
60	ATOM	717	CD1	PHE	89	-0.762	72.504	-24.067	1.00	31.56
	ATOM	718	CD2	PHE	89	-2.376	72.852	-22.314	1.00	30.96
	ATOM	719	CE1	PHE	89	-0.749	73.880	-24.316	1.00	33.76
	ATOM	720	CE2	PHE	89	-2.369	74.221	-22.553	1.00	27.38
	ATOM	721	CZ	PHE	89	-1.552	74.739	-23.551	1.00	30.29
65	ATOM	722	C	PHE	89	-1.223	68.836	-20.788	1.00	39.99
	ATOM	723	O	PHE	89	-1.375	68.466	-19.620	1.00	45.34

	ATOM	724	N	ARG	90	-0.871	67.977	-21.734	1.00	38.92
	ATOM	725	CA	ARG	90	-0.695	66.555	-21.386	1.00	41.66
	ATOM	726	CB	ARG	90	0.047	66.403	-20.042	1.00	39.24
	ATOM	727	CG	ARG	90	-0.435	65.232	-19.162	1.00	29.41
5	ATOM	728	CD	ARG	90	0.363	65.117	-17.850	1.00	31.13
	ATOM	729	NE	ARG	90	-0.262	64.196	-16.898	1.00	31.75
	ATOM	730	CZ	ARG	90	-0.796	64.550	-15.726	1.00	28.86
	ATOM	731	NH1	ARG	90	-0.788	65.810	-15.326	1.00	25.48
	ATOM	732	NH2	ARG	90	-1.373	63.639	-14.961	1.00	29.84
10	ATOM	733	C	ARG	90	-2.008	65.756	-21.289	1.00	39.23
	ATOM	734	O	ARG	90	-2.945	66.166	-20.607	1.00	31.33
	ATOM	735	N	CYS	91	-2.067	64.613	-21.977	1.00	40.12
	ATOM	736	CA	CYS	91	-3.246	63.762	-21.909	1.00	37.09
	ATOM	737	C	CYS	91	-3.001	62.288	-21.755	1.00	37.21
15	ATOM	738	O	CYS	91	-1.887	61.809	-21.927	1.00	37.46
	ATOM	739	CB	CYS	91	-4.171	63.968	-23.086	1.00	40.13
	ATOM	740	SG	CYS	91	-3.663	63.642	-24.812	1.00	41.64
	ATOM	741	N	ILE	92	-4.074	61.575	-21.430	1.00	34.02
	ATOM	742	CA	ILE	92	-4.009	60.139	-21.199	1.00	33.22
20	ATOM	743	CB	ILE	92	-4.060	59.844	-19.686	1.00	32.88
	ATOM	744	CG2	ILE	92	-3.799	58.374	-19.432	1.00	32.61
	ATOM	745	CG1	ILE	92	-3.011	60.690	-18.961	1.00	34.64
	ATOM	746	CD1	ILE	92	-3.095	60.631	-17.444	1.00	32.30
	ATOM	747	C	ILE	92	-5.144	59.377	-21.871	1.00	30.78
25	ATOM	748	O	ILE	92	-6.206	59.919	-22.100	1.00	33.43
	ATOM	749	N	GLN	93	-4.896	58.118	-22.209	1.00	29.35
	ATOM	750	CA	GLN	93	-5.899	57.260	-22.818	1.00	28.69
	ATOM	751	CB	GLN	93	-5.752	57.198	-24.339	1.00	28.17
	ATOM	752	CG	GLN	93	-6.411	58.314	-25.153	1.00	25.87
30	ATOM	753	CD	GLN	93	-7.912	58.396	-24.978	1.00	24.22
	ATOM	754	OE1	GLN	93	-8.400	59.074	-24.088	1.00	28.90
	ATOM	755	NE2	GLN	93	-8.651	57.699	-25.831	1.00	25.29
	ATOM	756	C	GLN	93	-5.684	55.865	-22.274	1.00	31.44
	ATOM	757	O	GLN	93	-4.578	55.348	-22.335	1.00	32.07
35	ATOM	758	N	ASN	94	-6.721	55.261	-21.713	1.00	35.77
	ATOM	759	CA	ASN	94	-6.589	53.901	-21.226	1.00	33.56
	ATOM	760	CB	ASN	94	-7.811	53.491	-20.402	1.00	34.70
	ATOM	761	CG	ASN	94	-7.747	53.983	-18.979	1.00	33.01
	ATOM	762	OD1	ASN	94	-6.764	53.769	-18.289	1.00	30.56
40	ATOM	763	ND2	ASN	94	-8.803	54.628	-18.526	1.00	36.61
	ATOM	764	C	ASN	94	-6.487	53.038	-22.484	1.00	35.80
	ATOM	765	O	ASN	94	-7.133	53.320	-23.495	1.00	36.63
	ATOM	766	N	MET	95	-5.668	51.996	-22.432	1.00	35.73
	ATOM	767	CA	MET	95	-5.493	51.124	-23.587	1.00	34.73
45	ATOM	768	CB	MET	95	-4.002	50.994	-23.912	1.00	32.97
	ATOM	769	CG	MET	95	-3.269	52.313	-24.110	1.00	27.35
	ATOM	770	SD	MET	95	-4.013	53.387	-25.355	1.00	25.14
	ATOM	771	CE	MET	95	-3.607	52.574	-26.863	1.00	15.55
	ATOM	772	C	MET	95	-6.093	49.734	-23.369	1.00	36.36
50	ATOM	773	O	MET	95	-6.193	49.264	-22.232	1.00	40.70
	ATOM	774	N	PRO	96	-6.526	49.071	-24.454	1.00	36.06
	ATOM	775	CD	PRO	96	-6.804	49.671	-25.759	1.00	35.32
	ATOM	776	CA	PRO	96	-7.107	47.729	-24.393	1.00	40.00
	ATOM	777	CB	PRO	96	-7.980	47.658	-25.641	1.00	33.97
55	ATOM	778	CG	PRO	96	-8.129	49.063	-26.068	1.00	37.05
	ATOM	779	C	PRO	96	-5.950	46.737	-24.484	1.00	46.34
	ATOM	780	O	PRO	96	-4.829	47.121	-24.814	1.00	47.59
	ATOM	781	N	GLU	97	-6.214	45.463	-24.212	1.00	52.14
	ATOM	782	CA	GLU	97	-5.166	44.452	-24.276	1.00	57.27
60	ATOM	783	CB	GLU	97	-5.498	43.280	-23.352	1.00	62.29
	ATOM	784	CG	GLU	97	-4.289	42.463	-22.906	1.00	72.31
	ATOM	785	CD	GLU	97	-3.819	42.845	-21.494	1.00	79.34
	ATOM	786	OE1	GLU	97	-4.619	42.672	-20.522	1.00	84.94
	ATOM	787	OE2	GLU	97	-2.657	43.320	-21.358	1.00	81.39
65	ATOM	788	C	GLU	97	-5.052	43.942	-25.708	1.00	56.67
	ATOM	789	O	GLU	97	-3.955	43.651	-26.193	1.00	59.18

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	ATOM	790	N	THR	98	-6.190	43.857	-26.386	1.00	57.55
	ATOM	791	CA	THR	98	-6.245	43.345	-27.750	1.00	59.71
	ATOM	792	CB	THR	98	-7.689	43.106	-28.172	1.00	59.24
	ATOM	793	OG1	THR	98	-8.407	44.351	-28.069	1.00	68.81
5	ATOM	794	CG2	THR	98	-8.333	42.029	-27.274	1.00	56.15
	ATOM	795	C	THR	98	-5.579	44.185	-28.839	1.00	60.32
	ATOM	796	O	THR	98	-4.367	44.070	-29.055	1.00	65.03
	ATOM	797	N	LEU	99	-6.365	45.007	-29.536	1.00	54.94
	ATOM	798	CA	LEU	99	-5.835	45.834	-30.624	1.00	51.13
10	ATOM	799	CB	LEU	99	-6.707	45.663	-31.866	1.00	50.56
	ATOM	800	CG	LEU	99	-6.580	44.336	-32.611	1.00	52.80
	ATOM	801	CD1	LEU	99	-7.779	44.131	-33.535	1.00	48.38
	ATOM	802	CD2	LEU	99	-5.277	44.330	-33.396	1.00	50.80
	ATOM	803	C	LEU	99	-5.722	47.322	-30.280	1.00	50.20
15	ATOM	804	O	LEU	99	-6.550	48.138	-30.712	1.00	50.50
	ATOM	805	N	PRO	100	-4.673	47.703	-29.529	1.00	47.97
	ATOM	806	CD	PRO	100	-3.524	46.878	-29.124	1.00	45.68
	ATOM	807	CA	PRO	100	-4.466	49.098	-29.130	1.00	46.54
	ATOM	808	CB	PRO	100	-3.100	49.067	-28.444	1.00	45.50
20	ATOM	809	CG	PRO	100	-2.985	47.659	-27.960	1.00	45.80
	ATOM	810	C	PRO	100	-4.485	50.060	-30.305	1.00	46.78
	ATOM	811	O	PRO	100	-3.714	49.918	-31.247	1.00	52.13
	ATOM	812	N	ASN	101	-5.371	51.040	-30.245	1.00	42.71
	ATOM	813	CA	ASN	101	-5.477	52.047	-31.282	1.00	41.37
25	ATOM	814	CB	ASN	101	-6.154	51.469	-32.519	1.00	43.24
	ATOM	815	CG	ASN	101	-5.186	50.735	-33.426	1.00	45.93
	ATOM	816	OD1	ASN	101	-4.329	51.347	-34.065	1.00	45.33
	ATOM	817	ND2	ASN	101	-5.314	49.411	-33.482	1.00	48.11
	ATOM	818	C	ASN	101	-6.292	53.202	-30.735	1.00	42.66
30	ATOM	819	O	ASN	101	-7.498	53.294	-30.975	1.00	44.79
	ATOM	820	N	ASN	102	-5.646	54.089	-29.984	1.00	40.46
	ATOM	821	CA	ASN	102	-6.384	55.202	-29.436	1.00	37.90
	ATOM	822	CB	ASN	102	-6.306	55.197	-27.917	1.00	34.01
	ATOM	823	CG	ASN	102	-7.387	54.336	-27.295	1.00	32.44
35	ATOM	824	OD1	ASN	102	-8.528	54.339	-27.750	1.00	29.07
	ATOM	825	ND2	ASN	102	-7.037	53.601	-26.250	1.00	33.79
	ATOM	826	C	ASN	102	-6.117	56.599	-29.964	1.00	39.20
	ATOM	827	O	ASN	102	-7.005	57.201	-30.555	1.00	46.07
	ATOM	828	N	SER	103	-4.927	57.137	-29.782	1.00	35.42
40	ATOM	829	CA	SER	103	-4.696	58.510	-30.264	1.00	40.28
	ATOM	830	CB	SER	103	-5.072	58.667	-31.758	1.00	40.68
	ATOM	831	OG	SER	103	-6.327	59.309	-31.947	1.00	31.76
	ATOM	832	C	SER	103	-5.493	59.528	-29.414	1.00	35.24
	ATOM	833	O	SER	103	-6.714	59.456	-29.287	1.00	24.65
45	ATOM	834	N	CYS	104	-4.764	60.457	-28.810	1.00	32.88
	ATOM	835	CA	CYS	104	-5.356	61.473	-27.978	1.00	31.77
	ATOM	836	C	CYS	104	-4.888	62.845	-28.495	1.00	30.58
	ATOM	837	O	CYS	104	-3.717	63.035	-28.813	1.00	29.23
	ATOM	838	CB	CYS	104	-4.951	61.265	-26.492	1.00	28.43
50	ATOM	839	SG	CYS	104	-5.387	62.756	-25.582	1.00	46.07
	ATOM	840	N	TYR	105	-5.820	63.788	-28.609	1.00	30.34
	ATOM	841	CA	TYR	105	-5.516	65.149	-29.048	1.00	26.12
	ATOM	842	CB	TYR	105	-6.466	65.581	-30.164	1.00	27.03
	ATOM	843	CG	TYR	105	-6.362	67.045	-30.561	1.00	25.61
55	ATOM	844	CD1	TYR	105	-5.607	67.446	-31.667	1.00	24.90
	ATOM	845	CE1	TYR	105	-5.511	68.782	-32.029	1.00	26.08
	ATOM	846	CD2	TYR	105	-7.017	68.031	-29.830	1.00	25.41
	ATOM	847	CE2	TYR	105	-6.922	69.370	-30.184	1.00	28.91
	ATOM	848	CZ	TYR	105	-6.169	69.739	-31.284	1.00	27.58
60	ATOM	849	OH	TYR	105	-6.088	71.067	-31.640	1.00	29.89
	ATOM	850	C	TYR	105	-5.695	66.086	-27.856	1.00	29.20
	ATOM	851	O	TYR	105	-6.582	65.891	-27.030	1.00	33.10
	ATOM	852	N	SER	106	-4.856	67.103	-27.760	1.00	30.48
	ATOM	853	CA	SER	106	-4.968	68.048	-26.661	1.00	29.22
65	ATOM	854	CB	SER	106	-4.263	67.495	-25.419	1.00	28.30
	ATOM	855	OG	SER	106	-4.395	68.365	-24.313	1.00	28.62

	ATOM	856	C	SER	106	-4.329	69.355	-27.098	1.00	29.94
	ATOM	857	O	SER	106	-3.332	69.341	-27.810	1.00	29.72
	ATOM	858	N	ALA	107	-4.914	70.479	-26.692	1.00	27.64
	ATOM	859	CA	ALA	107	-4.384	71.787	-27.051	1.00	24.33
5	ATOM	860	CB	ALA	107	-4.890	72.199	-28.422	1.00	21.30
	ATOM	861	C	ALA	107	-4.752	72.847	-26.025	1.00	25.56
	ATOM	862	O	ALA	107	-5.619	72.646	-25.185	1.00	28.03
	ATOM	863	N	GLY	108	-4.082	73.986	-26.102	1.00	26.07
	ATOM	864	CA	GLY	108	-4.360	75.070	-25.183	1.00	25.49
10	ATOM	865	C	GLY	108	-3.531	76.283	-25.538	1.00	29.03
	ATOM	866	O	GLY	108	-2.712	76.224	-26.444	1.00	32.20
	ATOM	867	N	ILE	109	-3.739	77.383	-24.828	1.00	29.34
	ATOM	868	CA	ILE	109	-2.997	78.608	-25.084	1.00	27.71
	ATOM	869	CB	ILE	109	-3.943	79.818	-25.161	1.00	27.48
15	ATOM	870	CG2	ILE	109	-3.154	81.087	-25.406	1.00	26.46
	ATOM	871	CG1	ILE	109	-4.958	79.598	-26.281	1.00	26.16
	ATOM	872	CD1	ILE	109	-6.038	80.640	-26.341	1.00	29.07
	ATOM	873	C	ILE	109	-1.988	78.837	-23.975	1.00	28.66
	ATOM	874	O	ILE	109	-2.261	78.555	-22.810	1.00	30.59
20	ATOM	875	N	ALA	110	-0.817	79.344	-24.335	1.00	25.60
	ATOM	876	CA	ALA	110	0.231	79.615	-23.358	1.00	26.36
	ATOM	877	CB	ALA	110	1.128	78.408	-23.212	1.00	20.26
	ATOM	878	C	ALA	110	1.054	80.811	-23.802	1.00	29.61
	ATOM	879	O	ALA	110	1.124	81.115	-24.983	1.00	27.61
25	ATOM	880	N	LYS	111	1.664	81.509	-22.855	1.00	33.77
	ATOM	881	CA	LYS	111	2.491	82.635	-23.222	1.00	35.09
	ATOM	882	CB	LYS	111	2.377	83.769	-22.210	1.00	39.38
	ATOM	883	CG	LYS	111	3.374	84.878	-22.521	1.00	49.74
	ATOM	884	CD	LYS	111	2.855	86.250	-22.191	1.00	53.95
30	ATOM	885	CE	LYS	111	3.830	87.308	-22.688	1.00	54.28
	ATOM	886	NZ	LYS	111	3.342	88.687	-22.344	1.00	60.85
	ATOM	887	C	LYS	111	3.934	82.153	-23.309	1.00	35.28
	ATOM	888	O	LYS	111	4.455	81.556	-22.374	1.00	36.58
	ATOM	889	N	LEU	112	4.565	82.409	-24.447	1.00	32.32
35	ATOM	890	CA	LEU	112	5.934	81.992	-24.680	1.00	29.67
	ATOM	891	CB	LEU	112	5.988	81.035	-25.869	1.00	27.04
	ATOM	892	CG	LEU	112	4.998	79.872	-25.845	1.00	24.38
	ATOM	893	CD1	LEU	112	5.080	79.102	-27.147	1.00	27.56
	ATOM	894	CD2	LEU	112	5.290	78.976	-24.673	1.00	19.52
40	ATOM	895	C	LEU	112	6.810	83.203	-24.963	1.00	32.58
	ATOM	896	O	LEU	112	6.309	84.291	-25.242	1.00	31.51
	ATOM	897	N	GLU	113	8.122	83.008	-24.898	1.00	35.75
	ATOM	898	CA	GLU	113	9.062	84.087	-25.145	1.00	40.08
	ATOM	899	CB	GLU	113	9.730	84.530	-23.856	1.00	44.23
45	ATOM	900	CG	GLU	113	8.814	84.755	-22.690	1.00	50.12
	ATOM	901	CD	GLU	113	9.466	85.657	-21.668	1.00	58.08
	ATOM	902	OE1	GLU	113	10.707	85.520	-21.452	1.00	61.13
	ATOM	903	OE2	GLU	113	8.738	86.502	-21.085	1.00	63.64
	ATOM	904	C	GLU	113	10.161	83.672	-26.090	1.00	41.63
50	ATOM	905	O	GLU	113	10.421	82.490	-26.271	1.00	48.29
	ATOM	906	N	GLU	114	10.813	84.669	-26.681	1.00	40.97
	ATOM	907	CA	GLU	114	11.932	84.449	-27.588	1.00	36.68
	ATOM	908	CB	GLU	114	12.689	85.742	-27.807	1.00	41.19
	ATOM	909	CG	GLU	114	12.341	86.520	-29.025	1.00	46.58
55	ATOM	910	CD	GLU	114	13.517	87.366	-29.461	1.00	51.12
	ATOM	911	OE1	GLU	114	14.543	86.773	-29.876	1.00	51.88
	ATOM	912	OE2	GLU	114	13.422	88.615	-29.373	1.00	57.08
	ATOM	913	C	GLU	114	12.907	83.493	-26.933	1.00	35.51
	ATOM	914	O	GLU	114	13.398	83.765	-25.844	1.00	38.92
60	ATOM	915	N	GLY	115	13.205	82.385	-27.594	1.00	32.99
	ATOM	916	CA	GLY	115	14.153	81.459	-27.017	1.00	31.79
	ATOM	917	C	GLY	115	13.521	80.259	-26.365	1.00	34.39
	ATOM	918	O	GLY	115	14.202	79.273	-26.116	1.00	39.20
	ATOM	919	N	ASP	116	12.233	80.342	-26.061	1.00	35.66
65	ATOM	920	CA	ASP	116	11.545	79.213	-25.459	1.00	38.09
	ATOM	921	CB	ASP	116	10.117	79.588	-25.042	1.00	36.00

	ATOM	922	CG	ASP	116	10.066	80.394	-23.763	1.00	35.09
	ATOM	923	OD1	ASP	116	11.073	80.433	-23.026	1.00	33.81
	ATOM	924	OD2	ASP	116	9.003	80.979	-23.484	1.00	35.24
	ATOM	925	C	ASP	116	11.476	78.112	-26.506	1.00	39.38
5	ATOM	926	O	ASP	116	11.490	78.385	-27.710	1.00	37.91
	ATOM	927	N	GLU	117	11.416	76.867	-26.053	1.00	41.54
	ATOM	928	CA	GLU	117	11.306	75.744	-26.975	1.00	37.92
	ATOM	929	CB	GLU	117	12.572	74.895	-26.955	1.00	38.90
	ATOM	930	CG	GLU	117	13.826	75.654	-27.305	1.00	48.15
10	ATOM	931	CD	GLU	117	15.049	74.762	-27.313	1.00	52.62
	ATOM	932	OE1	GLU	117	15.137	73.851	-26.444	1.00	55.49
	ATOM	933	OE2	GLU	117	15.925	74.980	-28.185	1.00	57.10
	ATOM	934	C	GLU	117	10.126	74.896	-26.543	1.00	34.02
	ATOM	935	O	GLU	117	9.854	74.770	-25.350	1.00	33.31
15	ATOM	936	N	LEU	118	9.412	74.343	-27.513	1.00	30.68
	ATOM	937	CA	LEU	118	8.277	73.481	-27.229	1.00	27.14
	ATOM	938	CB	LEU	118	7.046	73.938	-28.013	1.00	26.49
	ATOM	939	CG	LEU	118	6.435	75.294	-27.680	1.00	26.21
	ATOM	940	CD1	LEU	118	5.292	75.575	-28.628	1.00	23.05
20	ATOM	941	CD2	LEU	118	5.951	75.306	-26.245	1.00	25.38
	ATOM	942	C	LEU	118	8.634	72.064	-27.656	1.00	28.33
	ATOM	943	O	LEU	118	9.351	71.870	-28.640	1.00	31.46
	ATOM	944	N	GLN	119	8.154	71.070	-26.916	1.00	28.96
	ATOM	945	CA	GLN	119	8.414	69.683	-27.280	1.00	29.13
25	ATOM	946	CB	GLN	119	9.722	69.210	-26.662	1.00	28.23
	ATOM	947	CG	GLN	119	9.672	69.049	-25.173	1.00	36.38
	ATOM	948	CD	GLN	119	11.008	68.645	-24.597	1.00	36.57
	ATOM	949	OE1	GLN	119	11.085	68.194	-23.455	1.00	40.70
	ATOM	950	NE2	GLN	119	12.069	68.811	-25.377	1.00	36.80
30	ATOM	951	C	GLN	119	7.271	68.760	-26.870	1.00	29.43
	ATOM	952	O	GLN	119	6.529	69.050	-25.935	1.00	28.80
	ATOM	953	N	LEU	120	7.135	67.658	-27.600	1.00	32.26
	ATOM	954	CA	LEU	120	6.101	66.654	-27.366	1.00	29.96
	ATOM	955	CB	LEU	120	5.410	66.313	-28.694	1.00	31.17
35	ATOM	956	CG	LEU	120	4.110	65.512	-28.832	1.00	30.26
	ATOM	957	CD1	LEU	120	4.144	64.263	-27.987	1.00	27.10
	ATOM	958	CD2	LEU	120	2.956	66.391	-28.442	1.00	30.96
	ATOM	959	C	LEU	120	6.810	65.416	-26.812	1.00	30.51
	ATOM	960	O	LEU	120	7.596	64.773	-27.509	1.00	32.88
40	ATOM	961	N	ALA	121	6.525	65.081	-25.559	1.00	27.40
	ATOM	962	CA	ALA	121	7.156	63.935	-24.918	1.00	25.55
	ATOM	963	CB	ALA	121	7.994	64.415	-23.744	1.00	19.41
	ATOM	964	C	ALA	121	6.178	62.859	-24.445	1.00	29.89
	ATOM	965	O	ALA	121	5.114	63.160	-23.908	1.00	35.88
45	ATOM	966	N	ILE	122	6.551	61.601	-24.660	1.00	30.33
	ATOM	967	CA	ILE	122	5.739	60.471	-24.230	1.00	30.27
	ATOM	968	CB	ILE	122	5.579	59.433	-25.346	1.00	29.27
	ATOM	969	CG2	ILE	122	4.656	58.331	-24.879	1.00	30.38
	ATOM	970	CG1	ILE	122	5.013	60.101	-26.602	1.00	26.03
50	ATOM	971	CD1	ILE	122	4.841	59.173	-27.791	1.00	29.21
	ATOM	972	C	ILE	122	6.487	59.841	-23.060	1.00	33.20
	ATOM	973	O	ILE	122	7.588	59.321	-23.230	1.00	33.30
	ATOM	974	N	PRO	123	5.899	59.888	-21.852	1.00	34.70
	ATOM	975	CD	PRO	123	4.611	60.525	-21.558	1.00	34.40
55	ATOM	976	CA	PRO	123	6.484	59.341	-20.620	1.00	36.49
	ATOM	977	CB	PRO	123	5.610	59.940	-19.515	1.00	30.22
	ATOM	978	CG	PRO	123	4.862	61.056	-20.186	1.00	36.72
	ATOM	979	C	PRO	123	6.483	57.815	-20.549	1.00	39.64
	ATOM	980	O	PRO	123	5.933	57.239	-19.604	1.00	40.48
60	ATOM	981	N	ARG	124	7.093	57.166	-21.537	1.00	44.58
	ATOM	982	CA	ARG	124	7.154	55.712	-21.574	1.00	48.04
	ATOM	983	CB	ARG	124	5.938	55.149	-22.299	1.00	52.91
	ATOM	984	CG	ARG	124	4.612	55.618	-21.727	1.00	60.50
	ATOM	985	CD	ARG	124	3.946	54.460	-21.017	1.00	72.57
65	ATOM	986	NE	ARG	124	2.732	54.879	-20.295	1.00	84.90
	ATOM	987	CZ	ARG	124	2.689	55.281	-19.019	1.00	85.44

	ATOM	988	NH1	ARG	124	3.807	55.328	-18.293	1.00	87.69
	ATOM	989	NH2	ARG	124	1.522	55.641	-18.469	1.00	82.78
	ATOM	990	C	ARG	124	8.426	55.218	-22.243	1.00	48.39
	ATOM	991	O	ARG	124	8.946	55.861	-23.155	1.00	45.26
5	ATOM	992	N	GLU	125	8.905	54.066	-21.777	1.00	52.84
	ATOM	993	CA	GLU	125	10.129	53.442	-22.273	1.00	55.02
	ATOM	994	CB	GLU	125	10.285	52.053	-21.648	1.00	63.62
	ATOM	995	CG	GLU	125	8.959	51.373	-21.301	1.00	70.62
	ATOM	996	CD	GLU	125	8.080	52.229	-20.397	1.00	74.76
10	ATOM	997	OE1	GLU	125	8.484	52.482	-19.230	1.00	75.75
	ATOM	998	OE2	GLU	125	6.988	52.649	-20.856	1.00	80.71
	ATOM	999	C	GLU	125	10.241	53.342	-23.788	1.00	53.44
	ATOM	1000	O	GLU	125	11.061	54.032	-24.393	1.00	57.50
	ATOM	1001	N	ASN	126	9.464	52.463	-24.406	1.00	47.18
15	ATOM	1002	CA	ASN	126	9.508	52.356	-25.858	1.00	47.59
	ATOM	1003	CB	ASN	126	10.199	51.074	-26.312	1.00	51.03
	ATOM	1004	CG	ASN	126	11.632	51.318	-26.763	1.00	57.07
	ATOM	1005	OD1	ASN	126	12.542	51.513	-25.943	1.00	61.15
	ATOM	1006	ND2	ASN	126	11.838	51.329	-28.081	1.00	56.63
20	ATOM	1007	C	ASN	126	8.092	52.394	-26.363	1.00	47.06
	ATOM	1008	O	ASN	126	7.546	51.395	-26.840	1.00	45.61
	ATOM	1009	N	ALA	127	7.501	53.574	-26.236	1.00	43.50
	ATOM	1010	CA	ALA	127	6.130	53.800	-26.634	1.00	38.26
	ATOM	1011	CB	ALA	127	5.843	55.284	-26.629	1.00	37.97
25	ATOM	1012	C	ALA	127	5.819	53.217	-27.998	1.00	37.16
	ATOM	1013	O	ALA	127	6.529	53.464	-28.971	1.00	33.14
	ATOM	1014	N	GLN	128	4.761	52.417	-28.054	1.00	37.87
	ATOM	1015	CA	GLN	128	4.319	51.842	-29.316	1.00	37.19
	ATOM	1016	CB	GLN	128	3.599	50.522	-29.059	1.00	40.23
30	ATOM	1017	CG	GLN	128	4.541	49.448	-28.567	1.00	38.14
	ATOM	1018	CD	GLN	128	5.709	49.274	-29.509	1.00	39.51
	ATOM	1019	OE1	GLN	128	5.547	48.811	-30.643	1.00	40.36
	ATOM	1020	NE2	GLN	128	6.899	49.667	-29.056	1.00	41.53
	ATOM	1021	C	GLN	128	3.384	52.893	-29.910	1.00	35.55
35	ATOM	1022	O	GLN	128	2.216	53.016	-29.531	1.00	35.57
	ATOM	1023	N	ILE	129	3.936	53.653	-30.843	1.00	34.64
	ATOM	1024	CA	ILE	129	3.254	54.773	-31.475	1.00	29.60
	ATOM	1025	CB	ILE	129	4.017	56.083	-31.061	1.00	31.03
	ATOM	1026	CG2	ILE	129	4.287	56.986	-32.237	1.00	29.76
40	ATOM	1027	CG1	ILE	129	3.252	56.782	-29.960	1.00	26.37
	ATOM	1028	CD1	ILE	129	3.119	55.950	-28.739	1.00	34.70
	ATOM	1029	C	ILE	129	3.179	54.647	-32.994	1.00	29.55
	ATOM	1030	O	ILE	129	4.001	53.981	-33.606	1.00	33.42
	ATOM	1031	N	SER	130	2.184	55.281	-33.603	1.00	27.67
45	ATOM	1032	CA	SER	130	2.068	55.266	-35.058	1.00	25.00
	ATOM	1033	CB	SER	130	0.614	55.317	-35.495	1.00	22.09
	ATOM	1034	OG	SER	130	0.537	55.540	-36.896	1.00	20.89
	ATOM	1035	C	SER	130	2.778	56.500	-35.590	1.00	26.03
	ATOM	1036	O	SER	130	2.614	57.579	-35.047	1.00	30.77
50	ATOM	1037	N	LEU	131	3.559	56.353	-36.650	1.00	24.49
	ATOM	1038	CA	LEU	131	4.282	57.489	-37.201	1.00	28.07
	ATOM	1039	CB	LEU	131	5.761	57.142	-37.372	1.00	28.87
	ATOM	1040	CG	LEU	131	6.665	57.401	-36.167	1.00	30.76
	ATOM	1041	CD1	LEU	131	6.058	56.856	-34.908	1.00	28.98
55	ATOM	1042	CD2	LEU	131	8.007	56.768	-36.424	1.00	33.37
	ATOM	1043	C	LEU	131	3.730	58.018	-38.514	1.00	33.66
	ATOM	1044	O	LEU	131	4.479	58.537	-39.342	1.00	37.67
	ATOM	1045	N	ASP	132	2.422	57.888	-38.706	1.00	34.97
	ATOM	1046	CA	ASP	132	1.781	58.392	-39.916	1.00	34.31
60	ATOM	1047	CB	ASP	132	0.472	57.651	-40.190	1.00	42.68
	ATOM	1048	CG	ASP	132	0.695	56.261	-40.764	1.00	47.98
	ATOM	1049	OD1	ASP	132	-0.287	55.482	-40.833	1.00	50.41
	ATOM	1050	OD2	ASP	132	1.850	55.959	-41.155	1.00	46.42
	ATOM	1051	C	ASP	132	1.510	59.875	-39.731	1.00	34.32
65	ATOM	1052	O	ASP	132	1.003	60.298	-38.698	1.00	32.87
	ATOM	1053	N	GLY	133	1.857	60.658	-40.744	1.00	34.53

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	ATOM	1054	CA	GLY	133	1.678	62.094	-40.673	1.00	36.93
	ATOM	1055	C	GLY	133	0.291	62.606	-40.339	1.00	35.06
	ATOM	1056	O	GLY	133	0.145	63.735	-39.870	1.00	40.44
	ATOM	1057	N	ASP	134	-0.732	61.795	-40.566	1.00	31.08
5	ATOM	1058	CA	ASP	134	-2.082	62.244	-40.289	1.00	27.63
	ATOM	1059	CB	ASP	134	-3.040	61.718	-41.361	1.00	28.43
	ATOM	1060	CG	ASP	134	-3.079	60.207	-41.432	1.00	31.78
	ATOM	1061	OD1	ASP	134	-2.038	59.569	-41.193	1.00	35.68
	ATOM	1062	OD2	ASP	134	-4.154	59.655	-41.747	1.00	32.53
10	ATOM	1063	C	ASP	134	-2.579	61.906	-38.895	1.00	25.08
	ATOM	1064	O	ASP	134	-3.483	62.542	-38.402	1.00	19.71
	ATOM	1065	N	VAL	135	-1.970	60.935	-38.235	1.00	26.84
	ATOM	1066	CA	VAL	135	-2.430	60.588	-36.897	1.00	23.72
	ATOM	1067	CB	VAL	135	-2.528	59.072	-36.717	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-3.674	58.553	-37.540	1.00	22.76
	ATOM	1069	CG2	VAL	135	-1.230	58.408	-37.123	1.00	22.12
	ATOM	1070	C	VAL	135	-1.623	61.188	-35.751	1.00	26.54
	ATOM	1071	O	VAL	135	-2.150	61.350	-34.658	1.00	29.26
	ATOM	1072	N	THR	136	-0.351	61.511	-35.975	1.00	26.39
20	ATOM	1073	CA	THR	136	0.430	62.136	-34.909	1.00	28.40
	ATOM	1074	CB	THR	136	1.404	61.122	-34.219	1.00	28.21
	ATOM	1075	OG1	THR	136	2.738	61.330	-34.677	1.00	37.13
	ATOM	1076	CG2	THR	136	0.991	59.701	-34.509	1.00	26.90
	ATOM	1077	C	THR	136	1.172	63.373	-35.436	1.00	31.60
25	ATOM	1078	O	THR	136	2.024	63.283	-36.325	1.00	27.29
	ATOM	1079	N	PHE	137	0.813	64.533	-34.884	1.00	31.45
	ATOM	1080	CA	PHE	137	1.383	65.808	-35.302	1.00	26.64
	ATOM	1081	CB	PHE	137	0.497	66.405	-36.385	1.00	26.84
	ATOM	1082	CG	PHE	137	-0.971	66.287	-36.100	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-1.627	67.245	-35.333	1.00	27.77
	ATOM	1084	CD2	PHE	137	-1.704	65.225	-36.604	1.00	23.70
	ATOM	1085	CE1	PHE	137	-2.984	67.148	-35.076	1.00	27.32
	ATOM	1086	CE2	PHE	137	-3.063	65.120	-36.349	1.00	26.31
	ATOM	1087	CZ	PHE	137	-3.702	66.086	-35.585	1.00	29.43
35	ATOM	1088	C	PHE	137	1.564	66.778	-34.137	1.00	28.06
	ATOM	1089	O	PHE	137	1.010	66.569	-33.068	1.00	25.04
	ATOM	1090	N	PHE	138	2.337	67.842	-34.356	1.00	29.81
	ATOM	1091	CA	PHE	138	2.642	68.807	-33.302	1.00	29.93
	ATOM	1092	CB	PHE	138	4.093	68.587	-32.879	1.00	27.47
40	ATOM	1093	CG	PHE	138	4.481	69.288	-31.622	1.00	28.49
	ATOM	1094	CD1	PHE	138	3.526	69.650	-30.678	1.00	27.65
	ATOM	1095	CD2	PHE	138	5.819	69.584	-31.375	1.00	31.06
	ATOM	1096	CE1	PHE	138	3.900	70.300	-29.509	1.00	30.90
	ATOM	1097	CE2	PHE	138	6.207	70.233	-30.210	1.00	27.65
45	ATOM	1098	CZ	PHE	138	5.247	70.593	-29.275	1.00	27.41
	ATOM	1099	C	PHE	138	2.368	70.272	-33.694	1.00	30.91
	ATOM	1100	O	PHE	138	2.836	70.761	-34.716	1.00	24.94
	ATOM	1101	N	GLY	139	1.620	70.946	-32.817	1.00	36.94
	ATOM	1102	CA	GLY	139	1.130	72.314	-32.979	1.00	36.08
50	ATOM	1103	C	GLY	139	1.884	73.612	-33.133	1.00	36.48
	ATOM	1104	O	GLY	139	2.776	73.709	-33.955	1.00	43.61
	ATOM	1105	N	ALA	140	1.453	74.625	-32.379	1.00	36.60
	ATOM	1106	CA	ALA	140	2.028	75.984	-32.370	1.00	34.94
	ATOM	1107	CB	ALA	140	3.545	75.913	-32.391	1.00	36.94
55	ATOM	1108	C	ALA	140	1.550	76.986	-33.438	1.00	32.53
	ATOM	1109	O	ALA	140	1.940	76.923	-34.599	1.00	27.96
	ATOM	1110	N	LEU	141	0.716	77.930	-33.007	1.00	33.23
	ATOM	1111	CA	LEU	141	0.162	78.987	-33.864	1.00	36.11
	ATOM	1112	CB	LEU	141	-1.264	78.623	-34.276	1.00	37.53
60	ATOM	1113	CG	LEU	141	-2.120	79.589	-35.101	1.00	38.87
	ATOM	1114	CD1	LEU	141	-3.356	78.860	-35.570	1.00	40.67
	ATOM	1115	CD2	LEU	141	-2.529	80.802	-34.288	1.00	37.53
	ATOM	1116	C	LEU	141	0.152	80.296	-33.068	1.00	38.89
	ATOM	1117	O	LEU	141	-0.347	80.337	-31.946	1.00	39.25
65	ATOM	1118	N	LYS	142	0.689	81.372	-33.635	1.00	42.86
	ATOM	1119	CA	LYS	142	0.714	82.631	-32.899	1.00	42.96

	ATOM	1120	CB	LYS	142	1.895	83.485	-33.350	1.00	41.80
	ATOM	1121	CG	LYS	142	1.945	84.814	-32.634	1.00	46.88
	ATOM	1122	CD	LYS	142	3.312	85.461	-32.695	1.00	49.58
	ATOM	1123	CE	LYS	142	3.281	86.786	-31.951	1.00	51.07
5	ATOM	1124	NZ	LYS	142	4.612	87.439	-31.898	1.00	56.20
	ATOM	1125	C	LYS	142	-0.568	83.452	-32.978	1.00	43.41
	ATOM	1126	O	LYS	142	-1.089	83.707	-34.061	1.00	44.98
	ATOM	1127	N	LEU	143	-1.071	83.868	-31.820	1.00	42.81
	ATOM	1128	CA	LEU	143	-2.292	84.668	-31.750	1.00	40.65
10	ATOM	1129	CB	LEU	143	-2.967	84.488	-30.392	1.00	36.56
	ATOM	1130	CG	LEU	143	-3.353	83.075	-29.953	1.00	36.17
	ATOM	1131	CD1	LEU	143	-3.947	83.132	-28.558	1.00	37.95
	ATOM	1132	CD2	LEU	143	-4.344	82.480	-30.929	1.00	27.38
	ATOM	1133	C	LEU	143	-1.972	86.146	-31.938	1.00	43.14
15	ATOM	1134	O	LEU	143	-0.891	86.609	-31.559	1.00	46.28
	ATOM	1135	N	LEU	144	-2.905	86.889	-32.519	1.00	42.63
	ATOM	1136	CA	LEU	144	-2.693	88.315	-32.722	1.00	44.11
	ATOM	1137	CB	LEU	144	-3.628	88.849	-33.802	1.00	44.88
	ATOM	1138	CG	LEU	144	-3.373	88.280	-35.196	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-4.409	88.793	-36.159	1.00	47.71
	ATOM	1140	CD2	LEU	144	-1.982	88.669	-35.662	1.00	47.87
	ATOM	1141	C	LEU	144	-2.947	89.062	-31.428	1.00	45.66
	ATOM	1142	O	LEU	144	-3.637	88.509	-30.552	1.00	46.37
	ATOM	1143	OXT	LEU	144	-2.460	90.202	-31.312	1.00	50.66
25	END					50.903	67.374	64.558	0.00	0.00

TABLE 5

	11								
	ATOM	1	CB	VAL	1	-5.253	95.517	-32.692	1.00 69.28
5	ATOM	2	CG1	VAL	1	-3.761	95.624	-33.077	1.00 68.26
	ATOM	3	CG2	VAL	1	-6.008	96.806	-33.034	1.00 71.75
	ATOM	4	C	VAL	1	-5.393	93.010	-32.745	1.00 64.86
	ATOM	5	O	VAL	1	-4.543	92.304	-33.307	1.00 64.02
	ATOM	6	N	VAL	1	-7.402	94.367	-33.353	1.00 65.10
10	ATOM	7	CA	VAL	1	-5.910	94.298	-33.416	1.00 65.98
	ATOM	8	N	THR	2	-5.898	92.715	-31.545	1.00 61.42
	ATOM	9	CA	THR	2	-5.483	91.516	-30.819	1.00 56.64
	ATOM	10	CB	THR	2	-4.849	91.856	-29.459	1.00 56.77
	ATOM	11	OG1	THR	2	-5.765	92.653	-28.702	1.00 57.45
15	ATOM	12	CG2	THR	2	-3.540	92.615	-29.639	1.00 55.68
	ATOM	13	C	THR	2	-6.656	90.595	-30.543	1.00 53.08
	ATOM	14	O	THR	2	-7.810	90.965	-30.728	1.00 52.00
	ATOM	15	N	GLN	3	-6.347	89.391	-30.076	1.00 50.34
	ATOM	16	CA	GLN	3	-7.375	88.406	-29.766	1.00 45.97
20	ATOM	17	CB	GLN	3	-7.070	87.092	-30.476	1.00 46.09
	ATOM	18	CG	GLN	3	-6.752	87.249	-31.940	1.00 49.60
	ATOM	19	CD	GLN	3	-6.538	85.919	-32.616	1.00 50.43
	ATOM	20	OE1	GLN	3	-7.472	85.137	-32.759	1.00 50.94
	ATOM	21	NE2	GLN	3	-5.302	85.649	-33.030	1.00 49.99
25	ATOM	22	C	GLN	3	-7.454	88.158	-28.271	1.00 42.41
	ATOM	23	O	GLN	3	-6.558	87.565	-27.677	1.00 37.34
	ATOM	24	N	ASP	4	-8.535	88.619	-27.662	1.00 40.32
	ATOM	25	CA	ASP	4	-8.719	88.436	-26.234	1.00 39.37
	ATOM	26	CB	ASP	4	-9.926	89.244	-25.751	1.00 44.56
30	ATOM	27	CG	ASP	4	-9.715	90.738	-25.881	1.00 45.76
	ATOM	28	OD1	ASP	4	-8.821	91.140	-26.661	1.00 48.58
	ATOM	29	OD2	ASP	4	-10.443	91.505	-25.215	1.00 45.55
	ATOM	30	C	ASP	4	-8.946	86.967	-25.949	1.00 36.30
	ATOM	31	O	ASP	4	-9.529	86.250	-26.758	1.00 36.29
35	ATOM	32	N	CYS	5	-8.471	86.520	-24.797	1.00 35.11
	ATOM	33	CA	CYS	5	-8.656	85.139	-24.389	1.00 35.61
	ATOM	34	CB	CYS	5	-7.699	84.209	-25.139	1.00 34.96
	ATOM	35	SG	CYS	5	-6.031	84.811	-25.314	1.00 35.34
	ATOM	36	C	CYS	5	-8.462	85.021	-22.892	1.00 31.80
40	ATOM	37	O	CYS	5	-7.741	85.804	-22.286	1.00 31.85
	ATOM	38	N	LEU	6	-9.145	84.056	-22.298	1.00 27.72
	ATOM	39	CA	LEU	6	-9.054	83.814	-20.873	1.00 30.80
	ATOM	40	CB	LEU	6	-10.262	84.408	-20.158	1.00 29.25
	ATOM	41	CG	LEU	6	-10.376	84.184	-18.649	1.00 29.47
45	ATOM	42	CD1	LEU	6	-11.204	85.281	-18.035	1.00 30.65
	ATOM	43	CD2	LEU	6	-11.013	82.853	-18.380	1.00 35.27
	ATOM	44	C	LEU	6	-9.009	82.311	-20.678	1.00 31.83
	ATOM	45	O	LEU	6	-9.856	81.595	-21.197	1.00 33.45
	ATOM	46	N	GLN	7	-8.015	81.830	-19.940	1.00 32.23
50	ATOM	47	CA	GLN	7	-7.879	80.408	-19.711	1.00 30.26
	ATOM	48	CB	GLN	7	-6.552	79.921	-20.279	1.00 29.56
	ATOM	49	CG	GLN	7	-6.411	78.418	-20.373	1.00 25.09
	ATOM	50	CD	GLN	7	-5.189	78.019	-21.176	1.00 26.50
	ATOM	51	OE1	GLN	7	-4.052	78.182	-20.734	1.00 24.40
55	ATOM	52	NE2	GLN	7	-5.419	77.509	-22.373	1.00 27.03
	ATOM	53	C	GLN	7	-7.965	80.104	-18.231	1.00 32.11
	ATOM	54	O	GLN	7	-7.448	80.843	-17.404	1.00 34.24
	ATOM	55	N	LEU	8	-8.643	79.011	-17.913	1.00 33.73
	ATOM	56	CA	LEU	8	-8.821	78.572	-16.538	1.00 35.41
60	ATOM	57	CB	LEU	8	-10.321	78.461	-16.213	1.00 32.44
	ATOM	58	CG	LEU	8	-11.076	79.639	-15.579	1.00 27.69
	ATOM	59	CD1	LEU	8	-10.336	80.939	-15.760	1.00 26.43
	ATOM	60	CD2	LEU	8	-12.453	79.699	-16.181	1.00 18.64
	ATOM	61	C	LEU	8	-8.134	77.232	-16.324	1.00 36.87
65	ATOM	62	O	LEU	8	-8.061	76.398	-17.225	1.00 36.05
	ATOM	63	N	ILE	9	-7.638	77.036	-15.112	1.00 39.25

	ATOM	64	CA	ILE	9	-6.929	75.823	-14.738	1.00	35.57
	ATOM	65	CB	ILE	9	-5.442	76.176	-14.487	1.00	34.28
	ATOM	66	CG2	ILE	9	-4.925	75.548	-13.217	1.00	38.20
	ATOM	67	CG1	ILE	9	-4.613	75.748	-15.676	1.00	35.22
5	ATOM	68	CD1	ILE	9	-3.162	76.012	-15.456	1.00	43.63
	ATOM	69	C	ILE	9	-7.577	75.216	-13.490	1.00	35.28
	ATOM	70	O	ILE	9	-8.087	75.938	-12.639	1.00	33.75
	ATOM	71	N	ALA	10	-7.574	73.892	-13.386	1.00	35.78
	ATOM	72	CA	ALA	10	-8.162	73.249	-12.221	1.00	35.28
10	ATOM	73	CB	ALA	10	-8.141	71.746	-12.385	1.00	28.47
	ATOM	74	C	ALA	10	-7.409	73.643	-10.958	1.00	36.49
	ATOM	75	O	ALA	10	-6.184	73.725	-10.949	1.00	37.41
	ATOM	76	N	ASP	11	-8.152	73.890	-9.889	1.00	39.44
	ATOM	77	CA	ASP	11	-7.557	74.268	-8.612	1.00	40.77
15	ATOM	78	CB	ASP	11	-8.304	75.464	-8.021	1.00	38.90
	ATOM	79	CG	ASP	11	-7.817	75.821	-6.638	1.00	40.34
	ATOM	80	OD1	ASP	11	-6.644	75.517	-6.315	1.00	39.50
	ATOM	81	OD2	ASP	11	-8.608	76.414	-5.881	1.00	37.38
	ATOM	82	C	ASP	11	-7.599	73.089	-7.646	1.00	42.25
20	ATOM	83	O	ASP	11	-8.588	72.872	-6.945	1.00	40.54
	ATOM	84	N	SER	12	-6.506	72.337	-7.616	1.00	42.92
	ATOM	85	CA	SER	12	-6.392	71.146	-6.777	1.00	43.84
	ATOM	86	CB	SER	12	-5.078	70.434	-7.078	1.00	42.13
	ATOM	87	OG	SER	12	-3.982	71.303	-6.856	1.00	43.31
25	ATOM	88	C	SER	12	-6.481	71.397	-5.281	1.00	46.10
	ATOM	89	O	SER	12	-6.487	70.454	-4.489	1.00	43.90
	ATOM	90	N	GLU	13	-6.559	72.664	-4.892	1.00	46.72
	ATOM	91	CA	GLU	13	-6.631	72.998	-3.482	1.00	44.93
	ATOM	92	CB	GLU	13	-5.670	74.131	-3.172	1.00	46.71
30	ATOM	93	CG	GLU	13	-4.237	73.704	-3.334	1.00	58.25
	ATOM	94	CD	GLU	13	-3.291	74.544	-2.501	1.00	66.70
	ATOM	95	OE1	GLU	13	-3.485	74.577	-1.254	1.00	74.10
	ATOM	96	OE2	GLU	13	-2.361	75.166	-3.080	1.00	71.55
	ATOM	97	C	GLU	13	-8.020	73.311	-2.936	1.00	43.24
35	ATOM	98	O	GLU	13	-8.152	73.836	-1.828	1.00	44.25
	ATOM	99	N	THR	14	-9.054	73.007	-3.706	1.00	37.06
	ATOM	100	CA	THR	14	-10.414	73.206	-3.222	1.00	36.77
	ATOM	101	CB	THR	14	-11.041	74.550	-3.697	1.00	33.39
	ATOM	102	OG1	THR	14	-11.397	74.464	-5.076	1.00	41.69
40	ATOM	103	CG2	THR	14	-10.062	75.696	-3.501	1.00	31.66
	ATOM	104	C	THR	14	-11.230	72.025	-3.738	1.00	34.96
	ATOM	105	O	THR	14	-10.930	71.463	-4.784	1.00	35.04
	ATOM	106	N	PRO	15	-12.260	71.623	-2.994	1.00	37.11
	ATOM	107	CD	PRO	15	-12.685	72.206	-1.713	1.00	36.50
45	ATOM	108	CA	PRO	15	-13.122	70.495	-3.367	1.00	38.81
	ATOM	109	CB	PRO	15	-14.051	70.357	-2.161	1.00	39.15
	ATOM	110	CG	PRO	15	-13.292	71.020	-1.037	1.00	37.32
	ATOM	111	C	PRO	15	-13.911	70.748	-4.644	1.00	37.78
	ATOM	112	O	PRO	15	-14.216	71.893	-4.966	1.00	35.91
50	ATOM	113	N	THR	16	-14.249	69.688	-5.372	1.00	36.67
	ATOM	114	CA	THR	16	-15.028	69.868	-6.586	1.00	38.46
	ATOM	115	CB	THR	16	-15.168	68.559	-7.386	1.00	38.61
	ATOM	116	OG1	THR	16	-15.894	67.598	-6.613	1.00	42.44
	ATOM	117	CG2	THR	16	-13.796	67.999	-7.728	1.00	40.11
55	ATOM	118	C	THR	16	-16.404	70.305	-6.130	1.00	35.97
	ATOM	119	O	THR	16	-16.923	69.772	-5.158	1.00	39.85
	ATOM	120	N	ILE	17	-16.990	71.273	-6.821	1.00	33.66
	ATOM	121	CA	ILE	17	-18.307	71.770	-6.452	1.00	34.76
	ATOM	122	CB	ILE	17	-18.570	73.139	-7.094	1.00	32.44
60	ATOM	123	CG2	ILE	17	-19.988	73.588	-6.803	1.00	28.66
	ATOM	124	CG1	ILE	17	-17.544	74.152	-6.572	1.00	33.15
	ATOM	125	CD1	ILE	17	-17.650	75.529	-7.195	1.00	31.19
	ATOM	126	C	ILE	17	-19.440	70.826	-6.834	1.00	40.64
	ATOM	127	O	ILE	17	-19.547	70.402	-7.977	1.00	44.07
65	ATOM	128	N	GLN	18	-20.290	70.496	-5.865	1.00	45.91
	ATOM	129	CA	GLN	18	-21.425	69.610	-6.108	1.00	50.22

	ATOM	130	CB	GLN	18	-21.538	68.561	-5.007	1.00	53.09
	ATOM	131	CG	GLN	18	-21.501	67.153	-5.540	1.00	57.60
	ATOM	132	CD	GLN	18	-20.139	66.819	-6.092	1.00	61.39
	ATOM	133	OE1	GLN	18	-19.167	66.696	-5.342	1.00	68.36
5	ATOM	134	NE2	GLN	18	-20.050	66.690	-7.409	1.00	64.93
	ATOM	135	C	GLN	18	-22.720	70.401	-6.157	1.00	51.56
	ATOM	136	O	GLN	18	-22.985	71.229	-5.291	1.00	56.44
	ATOM	137	N	LYS	19	-23.533	70.141	-7.167	1.00	52.60
	ATOM	138	CA	LYS	19	-24.795	70.852	-7.287	1.00	56.10
10	ATOM	139	CB	LYS	19	-24.552	72.320	-7.639	1.00	55.55
	ATOM	140	CG	LYS	19	-25.847	73.063	-7.879	1.00	61.15
	ATOM	141	CD	LYS	19	-25.660	74.560	-8.012	1.00	64.42
	ATOM	142	CE	LYS	19	-27.017	75.244	-8.234	1.00	64.84
	ATOM	143	NZ	LYS	19	-26.906	76.734	-8.338	1.00	69.52
15	ATOM	144	C	LYS	19	-25.721	70.232	-8.322	1.00	57.41
	ATOM	145	O	LYS	19	-25.291	69.900	-9.429	1.00	61.02
	ATOM	146	N	GLY	20	-26.994	70.089	-7.955	1.00	58.84
	ATOM	147	CA	GLY	20	-27.980	69.514	-8.850	1.00	56.66
	ATOM	148	C	GLY	20	-27.519	68.217	-9.486	1.00	56.87
20	ATOM	149	O	GLY	20	-27.808	67.971	-10.658	1.00	58.64
	ATOM	150	N	SER	21	-26.810	67.391	-8.714	1.00	54.49
	ATOM	151	CA	SER	21	-26.295	66.105	-9.191	1.00	53.68
	ATOM	152	CB	SER	21	-27.452	65.187	-9.610	1.00	55.80
	ATOM	153	OG	SER	21	-27.862	65.433	-10.942	1.00	65.34
25	ATOM	154	C	SER	21	-25.285	66.262	-10.353	1.00	51.41
	ATOM	155	O	SER	21	-25.114	65.363	-11.194	1.00	47.74
	ATOM	156	N	TYR	22	-24.627	67.420	-10.385	1.00	46.25
	ATOM	157	CA	TYR	22	-23.614	67.734	-11.383	1.00	40.57
	ATOM	158	CB	TYR	22	-24.029	68.953	-12.210	1.00	43.27
30	ATOM	159	CG	TYR	22	-24.878	68.625	-13.411	1.00	49.18
	ATOM	160	CD1	TYR	22	-26.101	67.980	-13.270	1.00	52.61
	ATOM	161	CE1	TYR	22	-26.886	67.662	-14.383	1.00	51.92
	ATOM	162	CD2	TYR	22	-24.455	68.950	-14.695	1.00	50.01
	ATOM	163	CE2	TYR	22	-25.231	68.640	-15.813	1.00	50.94
35	ATOM	164	CZ	TYR	22	-26.443	67.996	-15.649	1.00	50.78
	ATOM	165	OH	TYR	22	-27.208	67.686	-16.752	1.00	53.87
	ATOM	166	C	TYR	22	-22.319	68.050	-10.643	1.00	37.85
	ATOM	167	O	TYR	22	-22.350	68.552	-9.520	1.00	38.70
	ATOM	168	N	THR	23	-21.182	67.742	-11.253	1.00	33.81
40	ATOM	169	CA	THR	23	-19.900	68.043	-10.626	1.00	32.40
	ATOM	170	CB	THR	23	-18.919	66.844	-10.663	1.00	31.58
	ATOM	171	OG1	THR	23	-19.512	65.702	-10.036	1.00	30.76
	ATOM	172	CG2	THR	23	-17.638	67.197	-9.937	1.00	24.08
	ATOM	173	C	THR	23	-19.266	69.188	-11.396	1.00	33.28
45	ATOM	174	O	THR	23	-19.213	69.159	-12.620	1.00	30.44
	ATOM	175	N	PHE	24	-18.792	70.194	-10.677	1.00	32.16
	ATOM	176	CA	PHE	24	-18.155	71.336	-11.308	1.00	30.68
	ATOM	177	CB	PHE	24	-18.921	72.620	-10.990	1.00	28.20
	ATOM	178	CG	PHE	24	-20.292	72.665	-11.586	1.00	31.00
50	ATOM	179	CD1	PHE	24	-21.364	72.071	-10.940	1.00	31.69
	ATOM	180	CD2	PHE	24	-20.510	73.276	-12.817	1.00	31.42
	ATOM	181	CE1	PHE	24	-22.630	72.084	-11.511	1.00	32.57
	ATOM	182	CE2	PHE	24	-21.769	73.294	-13.391	1.00	25.01
	ATOM	183	CZ	PHE	24	-22.831	72.696	-12.736	1.00	30.12
55	ATOM	184	C	PHE	24	-16.703	71.484	-10.872	1.00	31.73
	ATOM	185	O	PHE	24	-16.390	71.465	-9.686	1.00	33.31
	ATOM	186	N	VAL	25	-15.818	71.624	-11.845	1.00	32.92
	ATOM	187	CA	VAL	25	-14.408	71.787	-11.565	1.00	30.84
	ATOM	188	CB	VAL	25	-13.577	71.705	-12.856	1.00	30.29
60	ATOM	189	CG1	VAL	25	-12.116	71.918	-12.547	1.00	28.13
	ATOM	190	CG2	VAL	25	-13.789	70.372	-13.520	1.00	29.53
	ATOM	191	C	VAL	25	-14.156	73.149	-10.929	1.00	36.08
	ATOM	192	O	VAL	25	-14.714	74.157	-11.358	1.00	35.82
	ATOM	193	N	PRO	26	-13.329	73.193	-9.874	1.00	38.03
65	ATOM	194	CD	PRO	26	-12.861	72.033	-9.099	1.00	38.28
	ATOM	195	CA	PRO	26	-12.996	74.444	-9.179	1.00	38.08

	ATOM	196	CB	PRO	26	-12.426	73.964	-7.842	1.00	37.62
	ATOM	197	CG	PRO	26	-12.929	72.563	-7.703	1.00	36.31
	ATOM	198	C	PRO	26	-11.917	75.120	-10.023	1.00	40.85
	ATOM	199	O	PRO	26	-10.796	74.622	-10.109	1.00	42.27
5	ATOM	200	N	TRP	27	-12.234	76.244	-10.648	1.00	40.86
	ATOM	201	CA	TRP	27	-11.244	76.892	-11.500	1.00	37.05
	ATOM	202	CB	TRP	27	-11.927	77.583	-12.683	1.00	33.29
	ATOM	203	CG	TRP	27	-12.694	76.651	-13.551	1.00	31.06
	ATOM	204	CD2	TRP	27	-12.174	75.542	-14.286	1.00	27.79
10	ATOM	205	CE2	TRP	27	-13.260	74.930	-14.934	1.00	29.28
	ATOM	206	CE3	TRP	27	-10.898	75.006	-14.460	1.00	27.06
	ATOM	207	CD1	TRP	27	-14.030	76.669	-13.778	1.00	31.07
	ATOM	208	NE1	TRP	27	-14.382	75.641	-14.606	1.00	32.70
	ATOM	209	CZ2	TRP	27	-13.112	73.802	-15.746	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-10.750	73.884	-15.268	1.00	26.93
	ATOM	211	CH2	TRP	27	-11.853	73.295	-15.899	1.00	28.08
	ATOM	212	C	TRP	27	-10.333	77.883	-10.813	1.00	38.50
	ATOM	213	O	TRP	27	-10.649	78.419	-9.756	1.00	39.07
	ATOM	214	N	LEU	28	-9.187	78.105	-11.447	1.00	40.19
20	ATOM	215	CA	LEU	28	-8.178	79.043	-10.982	1.00	39.45
	ATOM	216	CB	LEU	28	-7.001	78.293	-10.369	1.00	44.18
	ATOM	217	CG	LEU	28	-6.060	79.128	-9.503	1.00	46.83
	ATOM	218	CD1	LEU	28	-6.810	79.568	-8.242	1.00	45.15
	ATOM	219	CD2	LEU	28	-4.821	78.304	-9.141	1.00	49.15
25	ATOM	220	C	LEU	28	-7.722	79.749	-12.252	1.00	39.80
	ATOM	221	O	LEU	28	-7.564	79.109	-13.282	1.00	42.40
	ATOM	222	N	LEU	29	-7.519	81.054	-12.197	1.00	33.58
	ATOM	223	CA	LEU	29	-7.094	81.759	-13.390	1.00	29.21
	ATOM	224	CB	LEU	29	-7.071	83.268	-13.154	1.00	27.16
30	ATOM	225	CG	LEU	29	-6.566	84.082	-14.349	1.00	22.36
	ATOM	226	CD1	LEU	29	-7.630	84.107	-15.434	1.00	21.47
	ATOM	227	CD2	LEU	29	-6.213	85.481	-13.920	1.00	14.53
	ATOM	228	C	LEU	29	-5.714	81.320	-13.867	1.00	29.98
	ATOM	229	O	LEU	29	-4.747	81.319	-13.112	1.00	30.36
35	ATOM	230	N	SER	30	-5.641	80.941	-15.136	1.00	29.88
	ATOM	231	CA	SER	30	-4.384	80.542	-15.740	1.00	28.49
	ATOM	232	CB	SER	30	-4.617	79.540	-16.870	1.00	26.13
	ATOM	233	OG	SER	30	-3.404	79.226	-17.517	1.00	20.25
	ATOM	234	C	SER	30	-3.803	81.838	-16.289	1.00	31.61
40	ATOM	235	O	SER	30	-2.666	82.197	-15.996	1.00	35.14
	ATOM	236	N	PHE	31	-4.601	82.544	-17.084	1.00	32.12
	ATOM	237	CA	PHE	31	-4.184	83.817	-17.647	1.00	30.76
	ATOM	238	CB	PHE	31	-3.054	83.620	-18.665	1.00	27.55
	ATOM	239	CG	PHE	31	-3.524	83.353	-20.068	1.00	31.23
45	ATOM	240	CD1	PHE	31	-3.893	84.401	-20.904	1.00	32.00
	ATOM	241	CD2	PHE	31	-3.611	82.050	-20.552	1.00	27.89
	ATOM	242	CE1	PHE	31	-4.338	84.157	-22.193	1.00	29.70
	ATOM	243	CE2	PHE	31	-4.055	81.801	-21.841	1.00	26.15
	ATOM	244	CZ	PHE	31	-4.420	82.854	-22.662	1.00	28.30
50	ATOM	245	C	PHE	31	-5.370	84.504	-18.302	1.00	32.06
	ATOM	246	O	PHE	31	-6.304	83.856	-18.774	1.00	32.26
	ATOM	247	N	LYS	32	-5.332	85.825	-18.309	1.00	34.83
	ATOM	248	CA	LYS	32	-6.387	86.610	-18.917	1.00	37.13
	ATOM	249	CB	LYS	32	-7.194	87.330	-17.839	1.00	36.55
55	ATOM	250	CG	LYS	32	-8.270	88.221	-18.394	1.00	40.62
	ATOM	251	CD	LYS	32	-8.857	89.125	-17.338	1.00	44.56
	ATOM	252	CE	LYS	32	-9.709	90.202	-18.014	1.00	46.29
	ATOM	253	NZ	LYS	32	-10.434	91.072	-17.042	1.00	50.58
	ATOM	254	C	LYS	32	-5.709	87.617	-19.842	1.00	36.39
60	ATOM	255	O	LYS	32	-4.728	88.257	-19.461	1.00	38.69
	ATOM	256	N	ARG	33	-6.218	87.743	-21.063	1.00	37.90
	ATOM	257	CA	ARG	33	-5.641	88.670	-22.026	1.00	35.74
	ATOM	258	CB	ARG	33	-4.857	87.883	-23.070	1.00	36.79
	ATOM	259	CG	ARG	33	-4.260	88.702	-24.191	1.00	31.09
65	ATOM	260	CD	ARG	33	-3.289	87.843	-24.992	1.00	31.62
	ATOM	261	NE	ARG	33	-2.549	88.592	-26.004	1.00	38.16

	ATOM	262	CZ	ARG	33	-2.849	88.597	-27.297	1.00	39.15
	ATOM	263	NH1	ARG	33	-3.876	87.888	-27.744	1.00	43.41
	ATOM	264	NH2	ARG	33	-2.128	89.311	-28.145	1.00	40.65
	ATOM	265	C	ARG	33	-6.743	89.487	-22.691	1.00	37.71
5	ATOM	266	O	ARG	33	-7.636	88.935	-23.340	1.00	41.16
	ATOM	267	N	GLY	34	-6.692	90.804	-22.526	1.00	36.59
	ATOM	268	CA	GLY	34	-7.708	91.640	-23.132	1.00	36.72
	ATOM	269	C	GLY	34	-8.870	91.916	-22.200	1.00	40.01
	ATOM	270	O	GLY	34	-8.793	91.671	-20.997	1.00	39.85
10	ATOM	271	N	SER	35	-9.962	92.404	-22.771	1.00	38.74
	ATOM	272	CA	SER	35	-11.142	92.755	-21.994	1.00	40.23
	ATOM	273	CB	SER	35	-11.420	94.241	-22.179	1.00	43.02
	ATOM	274	OG	SER	35	-11.507	94.558	-23.567	1.00	48.05
	ATOM	275	C	SER	35	-12.418	91.975	-22.323	1.00	40.54
15	ATOM	276	O	SER	35	-13.355	91.973	-21.526	1.00	43.74
	ATOM	277	N	ALA	36	-12.456	91.321	-23.481	1.00	35.56
	ATOM	278	CA	ALA	36	-13.638	90.583	-23.914	1.00	29.62
	ATOM	279	CB	ALA	36	-13.407	90.036	-25.308	1.00	25.47
	ATOM	280	C	ALA	36	-14.116	89.463	-22.994	1.00	31.23
20	ATOM	281	O	ALA	36	-15.291	89.116	-23.012	1.00	29.56
	ATOM	282	N	LEU	37	-13.220	88.897	-22.193	1.00	34.09
	ATOM	283	CA	LEU	37	-13.589	87.797	-21.306	1.00	32.32
	ATOM	284	CB	LEU	37	-13.005	86.490	-21.836	1.00	26.47
	ATOM	285	CG	LEU	37	-13.468	86.100	-23.237	1.00	25.72
25	ATOM	286	CD1	LEU	37	-12.519	85.097	-23.844	1.00	24.61
	ATOM	287	CD2	LEU	37	-14.874	85.553	-23.174	1.00	27.53
	ATOM	288	C	LEU	37	-13.134	88.010	-19.873	1.00	35.16
	ATOM	289	O	LEU	37	-12.055	88.533	-19.620	1.00	36.47
	ATOM	290	N	GLU	38	-13.971	87.589	-18.937	1.00	38.30
30	ATOM	291	CA	GLU	38	-13.673	87.720	-17.520	1.00	41.17
	ATOM	292	CB	GLU	38	-14.455	88.882	-16.918	1.00	46.15
	ATOM	293	CG	GLU	38	-13.776	90.229	-16.985	1.00	50.01
	ATOM	294	CD	GLU	38	-14.734	91.357	-16.623	1.00	51.18
	ATOM	295	OE1	GLU	38	-15.545	91.163	-15.680	1.00	48.40
35	ATOM	296	OE2	GLU	38	-14.673	92.430	-17.280	1.00	50.23
	ATOM	297	C	GLU	38	-14.057	86.464	-16.763	1.00	42.69
	ATOM	298	O	GLU	38	-14.897	85.690	-17.204	1.00	42.55
	ATOM	299	N	GLU	39	-13.440	86.260	-15.611	1.00	44.75
	ATOM	300	CA	GLU	39	-13.777	85.103	-14.799	1.00	46.50
40	ATOM	301	CB	GLU	39	-12.543	84.579	-14.071	1.00	47.52
	ATOM	302	CG	GLU	39	-11.639	85.685	-13.574	1.00	58.50
	ATOM	303	CD	GLU	39	-10.490	85.163	-12.733	1.00	62.35
	ATOM	304	OE1	GLU	39	-9.562	85.957	-12.426	1.00	64.58
	ATOM	305	OE2	GLU	39	-10.522	83.964	-12.370	1.00	63.04
45	ATOM	306	C	GLU	39	-14.794	85.635	-13.806	1.00	44.56
	ATOM	307	O	GLU	39	-14.636	86.731	-13.283	1.00	48.11
	ATOM	308	N	LYS	40	-15.850	84.875	-13.561	1.00	40.71
	ATOM	309	CA	LYS	40	-16.878	85.311	-12.637	1.00	38.08
	ATOM	310	CB	LYS	40	-17.920	86.154	-13.368	1.00	41.28
50	ATOM	311	CG	LYS	40	-19.136	86.510	-12.522	1.00	41.52
	ATOM	312	CD	LYS	40	-20.264	87.051	-13.389	1.00	42.73
	ATOM	313	CE	LYS	40	-21.543	87.264	-12.599	1.00	41.97
	ATOM	314	NZ	LYS	40	-22.663	87.676	-13.500	1.00	45.94
	ATOM	315	C	LYS	40	-17.562	84.128	-11.984	1.00	39.66
55	ATOM	316	O	LYS	40	-18.248	83.351	-12.643	1.00	36.18
	ATOM	317	N	GLU	41	-17.364	84.001	-10.679	1.00	40.16
	ATOM	318	CA	GLU	41	-17.973	82.930	-9.904	1.00	42.80
	ATOM	319	CB	GLU	41	-19.447	83.253	-9.692	1.00	45.31
	ATOM	320	CG	GLU	41	-19.642	84.624	-9.056	1.00	56.07
60	ATOM	321	CD	GLU	41	-21.058	85.170	-9.212	1.00	59.42
	ATOM	322	OE1	GLU	41	-21.555	85.256	-10.363	1.00	63.71
	ATOM	323	OE2	GLU	41	-21.673	85.521	-8.181	1.00	63.71
	ATOM	324	C	GLU	41	-17.805	81.569	-10.560	1.00	38.93
	ATOM	325	O	GLU	41	-18.776	80.852	-10.786	1.00	39.03
65	ATOM	326	N	ASN	42	-16.561	81.237	-10.877	1.00	34.69
	ATOM	327	CA	ASN	42	-16.218	79.961	-11.480	1.00	32.96

	ATOM	328	CB	ASN	42	-16.734	78.818	-10.613	1.00	32.65
	ATOM	329	CG	ASN	42	-15.816	77.636	-10.631	1.00	35.50
	ATOM	330	OD1	ASN	42	-16.257	76.499	-10.740	1.00	40.26
	ATOM	331	ND2	ASN	42	-14.523	77.895	-10.517	1.00	29.28
5	ATOM	332	C	ASN	42	-16.712	79.758	-12.904	1.00	31.44
	ATOM	333	O	ASN	42	-16.764	78.633	-13.392	1.00	22.78
	ATOM	334	N	LYS	43	-17.065	80.847	-13.569	1.00	33.26
	ATOM	335	CA	LYS	43	-17.550	80.776	-14.937	1.00	33.68
	ATOM	336	CB	LYS	43	-19.062	81.001	-14.973	1.00	33.55
10	ATOM	337	CG	LYS	43	-19.858	79.894	-14.329	1.00	40.49
	ATOM	338	CD	LYS	43	-21.329	80.225	-14.316	1.00	45.39
	ATOM	339	CE	LYS	43	-21.602	81.423	-13.429	1.00	52.57
	ATOM	340	NZ	LYS	43	-23.059	81.694	-13.322	1.00	62.06
	ATOM	341	C	LYS	43	-16.867	81.834	-15.772	1.00	32.05
15	ATOM	342	O	LYS	43	-16.240	82.738	-15.237	1.00	37.32
	ATOM	343	N	ILE	44	-16.971	81.719	-17.087	1.00	31.31
	ATOM	344	CA	ILE	44	-16.363	82.713	-17.946	1.00	29.52
	ATOM	345	CB	ILE	44	-15.676	82.069	-19.151	1.00	26.38
	ATOM	346	CG2	ILE	44	-15.096	83.146	-20.045	1.00	26.09
20	ATOM	347	CG1	ILE	44	-14.555	81.150	-18.666	1.00	28.61
	ATOM	348	CD1	ILE	44	-13.787	80.476	-19.762	1.00	21.51
	ATOM	349	C	ILE	44	-17.441	83.675	-18.414	1.00	31.60
	ATOM	350	O	ILE	44	-18.436	83.270	-19.011	1.00	34.50
	ATOM	351	N	LEU	45	-17.245	84.952	-18.115	1.00	31.44
25	ATOM	352	CA	LEU	45	-18.193	85.989	-18.493	1.00	31.40
	ATOM	353	CB	LEU	45	-18.342	86.989	-17.350	1.00	33.60
	ATOM	354	CG	LEU	45	-19.218	88.207	-17.634	1.00	34.36
	ATOM	355	CD1	LEU	45	-20.660	87.783	-17.819	1.00	29.00
	ATOM	356	CD2	LEU	45	-19.088	89.183	-16.480	1.00	32.42
30	ATOM	357	C	LEU	45	-17.771	86.725	-19.762	1.00	27.80
	ATOM	358	O	LEU	45	-16.641	87.188	-19.880	1.00	27.94
	ATOM	359	N	VAL	46	-18.695	86.824	-20.709	1.00	28.00
	ATOM	360	CA	VAL	46	-18.443	87.509	-21.968	1.00	31.53
	ATOM	361	CB	VAL	46	-19.328	86.915	-23.078	1.00	27.89
35	ATOM	362	CG1	VAL	46	-19.038	87.585	-24.399	1.00	26.10
	ATOM	363	CG2	VAL	46	-19.089	85.434	-23.173	1.00	23.39
	ATOM	364	C	VAL	46	-18.755	89.005	-21.809	1.00	33.94
	ATOM	365	O	VAL	46	-19.863	89.374	-21.414	1.00	39.68
	ATOM	366	N	LYS	47	-17.784	89.861	-22.118	1.00	33.59
40	ATOM	367	CA	LYS	47	-17.977	91.304	-21.991	1.00	37.05
	ATOM	368	CB	LYS	47	-16.812	91.928	-21.214	1.00	34.49
	ATOM	369	CG	LYS	47	-16.478	91.200	-19.924	1.00	37.53
	ATOM	370	CD	LYS	47	-17.682	91.121	-18.994	1.00	41.69
	ATOM	371	CE	LYS	47	-17.676	92.254	-17.986	1.00	42.95
45	ATOM	372	NZ	LYS	47	-17.403	93.557	-18.634	1.00	44.50
	ATOM	373	C	LYS	47	-18.140	92.017	-23.333	1.00	37.34
	ATOM	374	O	LYS	47	-18.520	93.183	-23.378	1.00	43.04
	ATOM	375	N	GLU	48	-17.848	91.314	-24.419	1.00	36.99
	ATOM	376	CA	GLU	48	-17.980	91.863	-25.765	1.00	37.31
50	ATOM	377	CB	GLU	48	-16.620	92.203	-26.352	1.00	36.53
	ATOM	378	CG	GLU	48	-15.914	93.360	-25.710	1.00	46.74
	ATOM	379	CD	GLU	48	-14.549	93.590	-26.333	1.00	52.39
	ATOM	380	OE1	GLU	48	-14.440	93.471	-27.585	1.00	51.33
	ATOM	381	OE2	GLU	48	-13.590	93.891	-25.576	1.00	54.55
55	ATOM	382	C	GLU	48	-18.594	90.783	-26.622	1.00	36.94
	ATOM	383	O	GLU	48	-18.095	89.666	-26.636	1.00	40.21
	ATOM	384	N	THR	49	-19.656	91.092	-27.354	1.00	36.34
	ATOM	385	CA	THR	49	-20.258	90.046	-28.165	1.00	35.48
	ATOM	386	CB	THR	49	-21.728	90.378	-28.548	1.00	31.16
60	ATOM	387	OG1	THR	49	-21.768	90.942	-29.857	1.00	32.48
	ATOM	388	CG2	THR	49	-22.329	91.356	-27.569	1.00	30.36
	ATOM	389	C	THR	49	-19.423	89.793	-29.423	1.00	34.59
	ATOM	390	O	THR	49	-18.745	90.688	-29.927	1.00	35.13
	ATOM	391	N	GLY	50	-19.463	88.554	-29.907	1.00	32.49
65	ATOM	392	CA	GLY	50	-18.713	88.195	-31.091	1.00	28.17
	ATOM	393	C	GLY	50	-18.590	86.693	-31.216	1.00	29.99

	ATOM	394	O	GLY	50	-19.345	85.958	-30.601	1.00	29.76
	ATOM	395	N	TYR	51	-17.638	86.240	-32.024	1.00	31.68
	ATOM	396	CA	TYR	51	-17.399	84.814	-32.218	1.00	32.88
	ATOM	397	CB	TYR	51	-17.169	84.501	-33.701	1.00	35.97
5	ATOM	398	CG	TYR	51	-18.414	84.637	-34.527	1.00	43.03
	ATOM	399	CD1	TYR	51	-18.848	85.889	-34.973	1.00	44.15
	ATOM	400	CE1	TYR	51	-20.054	86.032	-35.661	1.00	47.81
	ATOM	401	CD2	TYR	51	-19.211	83.526	-34.792	1.00	47.08
	ATOM	402	CE2	TYR	51	-20.420	83.652	-35.475	1.00	51.79
10	ATOM	403	CZ	TYR	51	-20.840	84.909	-35.905	1.00	51.15
	ATOM	404	OH	TYR	51	-22.063	85.039	-36.536	1.00	52.20
	ATOM	405	C	TYR	51	-16.200	84.350	-31.402	1.00	31.97
	ATOM	406	O	TYR	51	-15.113	84.914	-31.495	1.00	30.01
	ATOM	407	N	PHE	52	-16.405	83.314	-30.597	1.00	31.50
15	ATOM	408	CA	PHE	52	-15.335	82.797	-29.761	1.00	30.41
	ATOM	409	CB	PHE	52	-15.680	82.968	-28.280	1.00	31.30
	ATOM	410	CG	PHE	52	-15.917	84.385	-27.862	1.00	29.70
	ATOM	411	CD1	PHE	52	-17.079	85.046	-28.225	1.00	26.20
	ATOM	412	CD2	PHE	52	-14.978	85.056	-27.092	1.00	28.38
20	ATOM	413	CE1	PHE	52	-17.298	86.349	-27.828	1.00	27.17
	ATOM	414	CE2	PHE	52	-15.191	86.361	-26.691	1.00	27.76
	ATOM	415	CZ	PHE	52	-16.351	87.009	-27.058	1.00	27.27
	ATOM	416	C	PHE	52	-15.032	81.332	-29.999	1.00	27.65
	ATOM	417	O	PHE	52	-15.915	80.546	-30.317	1.00	24.85
25	ATOM	418	N	PHE	53	-13.764	80.984	-29.844	1.00	27.18
	ATOM	419	CA	PHE	53	-13.315	79.606	-29.960	1.00	27.05
	ATOM	420	CB	PHE	53	-11.913	79.536	-30.552	1.00	28.76
	ATOM	421	CG	PHE	53	-11.315	78.167	-30.520	1.00	28.00
	ATOM	422	CD1	PHE	53	-11.793	77.163	-31.348	1.00	26.49
30	ATOM	423	CD2	PHE	53	-10.280	77.871	-29.645	1.00	28.04
	ATOM	424	CE1	PHE	53	-11.247	75.885	-31.306	1.00	26.54
	ATOM	425	CE2	PHE	53	-9.732	76.593	-29.599	1.00	25.78
	ATOM	426	CZ	PHE	53	-10.218	75.602	-30.431	1.00	24.39
	ATOM	427	C	PHE	53	-13.275	79.159	-28.501	1.00	27.56
35	ATOM	428	O	PHE	53	-12.572	79.755	-27.691	1.00	25.39
	ATOM	429	N	ILE	54	-14.042	78.125	-28.168	1.00	28.87
	ATOM	430	CA	ILE	54	-14.117	77.635	-26.796	1.00	28.95
	ATOM	431	CB	ILE	54	-15.578	77.679	-26.305	1.00	29.60
	ATOM	432	CG2	ILE	54	-15.631	77.442	-24.805	1.00	27.42
40	ATOM	433	CG1	ILE	54	-16.187	79.043	-26.640	1.00	26.41
	ATOM	434	CD1	ILE	54	-17.678	79.108	-26.520	1.00	28.70
	ATOM	435	C	ILE	54	-13.574	76.211	-26.680	1.00	28.57
	ATOM	436	O	ILE	54	-13.856	75.358	-27.514	1.00	29.17
	ATOM	437	N	TYR	55	-12.795	75.948	-25.638	1.00	26.97
45	ATOM	438	CA	TYR	55	-12.212	74.622	-25.472	1.00	24.22
	ATOM	439	CB	TYR	55	-10.795	74.612	-26.047	1.00	21.82
	ATOM	440	CG	TYR	55	-9.862	75.616	-25.416	1.00	27.68
	ATOM	441	CD1	TYR	55	-9.088	75.279	-24.309	1.00	27.01
	ATOM	442	CE1	TYR	55	-8.249	76.207	-23.719	1.00	27.44
50	ATOM	443	CD2	TYR	55	-9.767	76.911	-25.914	1.00	27.16
	ATOM	444	CE2	TYR	55	-8.931	77.846	-25.325	1.00	25.39
	ATOM	445	CZ	TYR	55	-8.178	77.490	-24.231	1.00	29.02
	ATOM	446	OH	TYR	55	-7.360	78.420	-23.644	1.00	31.82
	ATOM	447	C	TYR	55	-12.192	74.156	-24.032	1.00	25.16
55	ATOM	448	O	TYR	55	-12.345	74.940	-23.124	1.00	28.47
	ATOM	449	N	GLY	56	-12.014	72.861	-23.829	1.00	27.44
	ATOM	450	CA	GLY	56	-11.983	72.336	-22.481	1.00	22.07
	ATOM	451	C	GLY	56	-11.514	70.898	-22.438	1.00	25.83
	ATOM	452	O	GLY	56	-11.772	70.128	-23.358	1.00	26.00
60	ATOM	453	N	GLN	57	-10.802	70.546	-21.375	1.00	22.56
	ATOM	454	CA	GLN	57	-10.308	69.192	-21.185	1.00	20.91
	ATOM	455	CB	GLN	57	-8.857	69.046	-21.644	1.00	17.87
	ATOM	456	CG	GLN	57	-8.278	67.672	-21.331	1.00	16.86
	ATOM	457	CD	GLN	57	-6.870	67.470	-21.849	1.00	22.06
65	ATOM	458	OE1	GLN	57	-6.634	67.474	-23.052	1.00	27.26
	ATOM	459	NE2	GLN	57	-5.924	67.285	-20.936	1.00	20.20

	ATOM	460	C	GLN	57	-10.400	68.836	-19.716	1.00	24.23
	ATOM	461	O	GLN	57	-10.200	69.690	-18.854	1.00	25.16
	ATOM	462	N	VAL	58	-10.705	67.571	-19.446	1.00	24.62
	ATOM	463	CA	VAL	58	-10.828	67.052	-18.088	1.00	26.68
5	ATOM	464	CB	VAL	58	-12.311	66.931	-17.661	1.00	25.55
	ATOM	465	CG1	VAL	58	-12.415	66.202	-16.344	1.00	15.69
	ATOM	466	CG2	VAL	58	-12.942	68.302	-17.546	1.00	25.77
	ATOM	467	C	VAL	58	-10.226	65.662	-18.070	1.00	31.06
	ATOM	468	O	VAL	58	-10.410	64.913	-19.020	1.00	27.74
10	ATOM	469	N	LEU	59	-9.506	65.319	-17.002	1.00	33.82
	ATOM	470	CA	LEU	59	-8.920	63.981	-16.864	1.00	29.85
	ATOM	471	CB	LEU	59	-7.527	64.043	-16.246	1.00	29.62
	ATOM	472	CG	LEU	59	-6.580	62.866	-16.522	1.00	28.11
	ATOM	473	CD1	LEU	59	-5.411	62.933	-15.561	1.00	26.67
15	ATOM	474	CD2	LEU	59	-7.282	61.549	-16.377	1.00	22.45
	ATOM	475	C	LEU	59	-9.828	63.168	-15.945	1.00	30.98
	ATOM	476	O	LEU	59	-9.968	63.478	-14.764	1.00	32.52
	ATOM	477	N	TYR	60	-10.448	62.127	-16.493	1.00	33.01
	ATOM	478	CA	TYR	60	-11.339	61.284	-15.709	1.00	34.51
20	ATOM	479	CB	TYR	60	-12.496	60.798	-16.576	1.00	40.10
	ATOM	480	CG	TYR	60	-13.338	61.923	-17.092	1.00	45.37
	ATOM	481	CD1	TYR	60	-13.266	62.322	-18.427	1.00	47.13
	ATOM	482	CE1	TYR	60	-13.984	63.420	-18.881	1.00	50.28
	ATOM	483	CD2	TYR	60	-14.152	62.644	-16.229	1.00	44.35
25	ATOM	484	CE2	TYR	60	-14.872	63.738	-16.665	1.00	48.81
	ATOM	485	CZ	TYR	60	-14.781	64.125	-17.986	1.00	51.91
	ATOM	486	OH	TYR	60	-15.467	65.241	-18.398	1.00	53.77
	ATOM	487	C	TYR	60	-10.639	60.091	-15.082	1.00	35.49
	ATOM	488	O	TYR	60	-10.031	59.276	-15.768	1.00	36.13
30	ATOM	489	N	THR	61	-10.731	60.002	-13.763	1.00	36.19
	ATOM	490	CA	THR	61	-10.131	58.908	-13.024	1.00	35.36
	ATOM	491	CB	THR	61	-9.080	59.427	-12.040	1.00	34.22
	ATOM	492	OG1	THR	61	-9.658	60.447	-11.221	1.00	34.22
	ATOM	493	CG2	THR	61	-7.888	59.993	-12.788	1.00	25.45
35	ATOM	494	C	THR	61	-11.258	58.219	-12.268	1.00	37.82
	ATOM	495	O	THR	61	-11.076	57.703	-11.169	1.00	43.70
	ATOM	496	N	ASP	62	-12.431	58.229	-12.886	1.00	37.64
	ATOM	497	CA	ASP	62	-13.641	57.640	-12.343	1.00	38.08
	ATOM	498	CB	ASP	62	-14.761	58.667	-12.472	1.00	38.42
40	ATOM	499	CG	ASP	62	-16.026	58.249	-11.779	1.00	41.25
	ATOM	500	OD1	ASP	62	-16.680	59.130	-11.173	1.00	38.28
	ATOM	501	OD2	ASP	62	-16.365	57.050	-11.855	1.00	41.37
	ATOM	502	C	ASP	62	-13.946	56.393	-13.168	1.00	40.90
	ATOM	503	O	ASP	62	-13.755	56.404	-14.376	1.00	46.73
45	ATOM	504	N	LYS	63	-14.414	55.316	-12.541	1.00	42.62
	ATOM	505	CA	LYS	63	-14.692	54.100	-13.305	1.00	40.92
	ATOM	506	CB	LYS	63	-14.187	52.870	-12.550	1.00	42.79
	ATOM	507	CG	LYS	63	-14.786	52.674	-11.174	1.00	45.91
	ATOM	508	CD	LYS	63	-14.058	51.554	-10.432	1.00	46.22
50	ATOM	509	CE	LYS	63	-12.572	51.904	-10.223	1.00	49.97
	ATOM	510	NZ	LYS	63	-11.743	50.762	-9.702	1.00	49.98
	ATOM	511	C	LYS	63	-16.150	53.893	-13.701	1.00	42.59
	ATOM	512	O	LYS	63	-16.552	52.785	-14.032	1.00	42.46
	ATOM	513	N	THR	64	-16.923	54.972	-13.707	1.00	44.06
55	ATOM	514	CA	THR	64	-18.336	54.918	-14.055	1.00	44.94
	ATOM	515	CB	THR	64	-19.028	56.254	-13.683	1.00	43.79
	ATOM	516	OG1	THR	64	-18.797	56.536	-12.300	1.00	46.20
	ATOM	517	CG2	THR	64	-20.533	56.176	-13.913	1.00	42.80
	ATOM	518	C	THR	64	-18.668	54.576	-15.517	1.00	45.82
60	ATOM	519	O	THR	64	-19.526	55.200	-16.114	1.00	53.10
	ATOM	520	N	TYR	65	-18.002	53.584	-16.088	1.00	45.11
	ATOM	521	CA	TYR	65	-18.256	53.136	-17.465	1.00	45.72
	ATOM	522	CB	TYR	65	-19.555	52.309	-17.519	1.00	40.46
	ATOM	523	CG	TYR	65	-20.809	53.071	-17.886	1.00	41.90
65	ATOM	524	CD1	TYR	65	-21.181	53.249	-19.220	1.00	42.47
	ATOM	525	CE1	TYR	65	-22.341	53.965	-19.565	1.00	41.83

	ATOM	526	CD2	TYR	65	-21.628	53.626	-16.897	1.00	41.65
	ATOM	527	CE2	TYR	65	-22.788	54.346	-17.227	1.00	40.28
	ATOM	528	CZ	TYR	65	-23.135	54.510	-18.562	1.00	41.90
	ATOM	529	OH	TYR	65	-24.267	55.212	-18.891	1.00	43.84
5	ATOM	530	C	TYR	65	-18.276	54.198	-18.576	1.00	45.31
	ATOM	531	O	TYR	65	-17.924	53.908	-19.728	1.00	49.74
	ATOM	532	N	ALA	66	-18.690	55.414	-18.248	1.00	42.32
	ATOM	533	CA	ALA	66	-18.748	56.492	-19.226	1.00	39.57
	ATOM	534	CB	ALA	66	-20.006	56.378	-20.058	1.00	35.36
10	ATOM	535	C	ALA	66	-18.712	57.833	-18.514	1.00	39.85
	ATOM	536	O	ALA	66	-19.563	58.125	-17.672	1.00	37.60
	ATOM	537	N	MET	67	-17.711	58.639	-18.843	1.00	41.88
	ATOM	538	CA	MET	67	-17.565	59.955	-18.239	1.00	41.48
	ATOM	539	CB	MET	67	-16.292	60.028	-17.386	1.00	39.98
15	ATOM	540	CG	MET	67	-16.320	59.183	-16.119	1.00	37.31
	ATOM	541	SD	MET	67	-17.661	59.625	-15.006	1.00	35.46
	ATOM	542	CE	MET	67	-16.984	61.062	-14.172	1.00	34.06
	ATOM	543	C	MET	67	-17.512	61.004	-19.336	1.00	39.50
	ATOM	544	O	MET	67	-17.338	60.681	-20.512	1.00	38.44
20	ATOM	545	N	GLY	68	-17.665	62.262	-18.943	1.00	39.84
	ATOM	546	CA	GLY	68	-17.629	63.342	-19.908	1.00	37.61
	ATOM	547	C	GLY	68	-18.057	64.658	-19.305	1.00	33.76
	ATOM	548	O	GLY	68	-18.506	64.704	-18.165	1.00	35.58
	ATOM	549	N	HIS	69	-17.904	65.735	-20.063	1.00	35.07
25	ATOM	550	CA	HIS	69	-18.293	67.052	-19.585	1.00	33.70
	ATOM	551	CB	HIS	69	-17.065	67.852	-19.113	1.00	37.02
	ATOM	552	CG	HIS	69	-15.963	67.951	-20.123	1.00	34.65
	ATOM	553	CD2	HIS	69	-15.559	68.982	-20.899	1.00	35.63
	ATOM	554	ND1	HIS	69	-15.117	66.902	-20.412	1.00	37.68
30	ATOM	555	CE1	HIS	69	-14.239	67.282	-21.322	1.00	32.64
	ATOM	556	NE2	HIS	69	-14.485	68.541	-21.634	1.00	35.90
	ATOM	557	C	HIS	69	-19.048	67.833	-20.652	1.00	33.98
	ATOM	558	O	HIS	69	-19.101	67.431	-21.803	1.00	30.94
	ATOM	559	N	LEU	70	-19.636	68.953	-20.254	1.00	35.35
35	ATOM	560	CA	LEU	70	-20.389	69.794	-21.172	1.00	30.76
	ATOM	561	CB	LEU	70	-21.864	69.810	-20.787	1.00	29.59
	ATOM	562	CG	LEU	70	-22.524	68.515	-20.324	1.00	30.71
	ATOM	563	CD1	LEU	70	-23.888	68.830	-19.758	1.00	25.60
	ATOM	564	CD2	LEU	70	-22.620	67.543	-21.475	1.00	27.39
40	ATOM	565	C	LEU	70	-19.885	71.233	-21.100	1.00	33.19
	ATOM	566	O	LEU	70	-19.604	71.744	-20.016	1.00	37.07
	ATOM	567	N	ILE	71	-19.744	71.881	-22.250	1.00	30.93
	ATOM	568	CA	ILE	71	-19.353	73.281	-22.255	1.00	27.94
	ATOM	569	CB	ILE	71	-18.372	73.610	-23.383	1.00	28.22
45	ATOM	570	CG2	ILE	71	-18.264	75.110	-23.561	1.00	24.54
	ATOM	571	CG1	ILE	71	-17.000	73.039	-23.046	1.00	26.34
	ATOM	572	CD1	ILE	71	-16.012	73.133	-24.175	1.00	37.50
	ATOM	573	C	ILE	71	-20.696	73.944	-22.514	1.00	30.32
	ATOM	574	O	ILE	71	-21.287	73.745	-23.570	1.00	23.07
50	ATOM	575	N	GLN	72	-21.193	74.704	-21.542	1.00	29.74
	ATOM	576	CA	GLN	72	-22.498	75.329	-21.699	1.00	27.85
	ATOM	577	CB	GLN	72	-23.407	74.876	-20.567	1.00	26.70
	ATOM	578	CG	GLN	72	-23.414	73.384	-20.397	1.00	28.28
	ATOM	579	CD	GLN	72	-24.456	72.924	-19.417	1.00	33.11
55	ATOM	580	OE1	GLN	72	-24.500	73.395	-18.284	1.00	41.22
	ATOM	581	NE2	GLN	72	-25.302	71.996	-19.842	1.00	30.18
	ATOM	582	C	GLN	72	-22.501	76.841	-21.773	1.00	29.85
	ATOM	583	O	GLN	72	-21.572	77.506	-21.311	1.00	28.20
	ATOM	584	N	ARG	73	-23.574	77.371	-22.349	1.00	30.90
60	ATOM	585	CA	ARG	73	-23.747	78.806	-22.499	1.00	30.79
	ATOM	586	CB	ARG	73	-23.763	79.161	-23.980	1.00	29.18
	ATOM	587	CG	ARG	73	-24.026	80.610	-24.256	1.00	28.66
	ATOM	588	CD	ARG	73	-24.573	80.781	-25.642	1.00	29.25
	ATOM	589	NE	ARG	73	-24.792	82.182	-25.964	1.00	35.78
65	ATOM	590	CZ	ARG	73	-25.572	82.606	-26.947	1.00	35.27
	ATOM	591	NH1	ARG	73	-26.218	81.733	-27.706	1.00	34.95

	ATOM	592	NH2	ARG	73	-25.699	83.902	-27.170	1.00	33.33
	ATOM	593	C	ARG	73	-25.044	79.294	-21.856	1.00	32.42
	ATOM	594	O	ARG	73	-26.110	78.750	-22.122	1.00	34.19
	ATOM	595	N	LYS	74	-24.941	80.304	-20.999	1.00	35.90
5	ATOM	596	CA	LYS	74	-26.116	80.895	-20.359	1.00	38.57
	ATOM	597	CB	LYS	74	-25.848	81.218	-18.886	1.00	43.52
	ATOM	598	CG	LYS	74	-25.781	80.008	-17.970	1.00	52.89
	ATOM	599	CD	LYS	74	-25.462	80.400	-16.508	1.00	58.14
	ATOM	600	CE	LYS	74	-25.386	79.160	-15.614	1.00	56.75
10	ATOM	601	NZ	LYS	74	-25.036	79.501	-14.203	1.00	63.98
	ATOM	602	C	LYS	74	-26.417	82.197	-21.092	1.00	40.11
	ATOM	603	O	LYS	74	-25.698	83.190	-20.924	1.00	37.55
	ATOM	604	N	LYS	75	-27.472	82.192	-21.899	1.00	34.48
	ATOM	605	CA	LYS	75	-27.856	83.373	-22.664	1.00	34.24
15	ATOM	606	CB	LYS	75	-28.969	83.015	-23.652	1.00	36.16
	ATOM	607	CG	LYS	75	-28.667	81.894	-24.645	1.00	35.99
	ATOM	608	CD	LYS	75	-29.891	81.660	-25.541	1.00	40.91
	ATOM	609	CE	LYS	75	-29.645	80.570	-26.575	1.00	46.16
	ATOM	610	NZ	LYS	75	-30.831	80.256	-27.447	1.00	46.03
20	ATOM	611	C	LYS	75	-28.354	84.498	-21.752	1.00	30.92
	ATOM	612	O	LYS	75	-29.062	84.244	-20.796	1.00	27.47
	ATOM	613	N	VAL	76	-27.982	85.739	-22.059	1.00	32.82
	ATOM	614	CA	VAL	76	-28.427	86.898	-21.280	1.00	31.51
	ATOM	615	CB	VAL	76	-27.600	88.166	-21.560	1.00	29.19
25	ATOM	616	CG1	VAL	76	-27.473	88.977	-20.302	1.00	27.09
	ATOM	617	CG2	VAL	76	-26.267	87.815	-22.119	1.00	34.94
	ATOM	618	C	VAL	76	-29.834	87.234	-21.728	1.00	33.40
	ATOM	619	O	VAL	76	-30.692	87.578	-20.922	1.00	32.92
	ATOM	620	N	HIS	77	-30.047	87.149	-23.037	1.00	34.88
30	ATOM	621	CA	HIS	77	-31.333	87.441	-23.645	1.00	34.59
	ATOM	622	CB	HIS	77	-31.146	88.367	-24.851	1.00	35.92
	ATOM	623	CG	HIS	77	-30.469	89.662	-24.522	1.00	34.17
	ATOM	624	CD2	HIS	77	-29.743	90.506	-25.288	1.00	31.44
	ATOM	625	ND1	HIS	77	-30.528	90.238	-23.270	1.00	35.05
35	ATOM	626	CE1	HIS	77	-29.867	91.380	-23.276	1.00	29.84
	ATOM	627	NE2	HIS	77	-29.383	91.566	-24.490	1.00	38.35
	ATOM	628	C	HIS	77	-31.991	86.139	-24.068	1.00	35.63
	ATOM	629	O	HIS	77	-31.325	85.246	-24.593	1.00	40.24
	ATOM	630	N	VAL	78	-33.304	86.038	-23.865	1.00	37.37
40	ATOM	631	CA	VAL	78	-34.001	84.807	-24.189	1.00	41.32
	ATOM	632	CB	VAL	78	-34.580	84.185	-22.903	1.00	43.10
	ATOM	633	CG1	VAL	78	-35.182	82.834	-23.207	1.00	45.01
	ATOM	634	CG2	VAL	78	-33.477	84.011	-21.875	1.00	44.98
	ATOM	635	C	VAL	78	-35.064	84.785	-25.295	1.00	43.02
45	ATOM	636	O	VAL	78	-34.849	84.140	-26.319	1.00	49.79
	ATOM	637	N	PHE	79	-36.202	85.447	-25.125	1.00	41.33
	ATOM	638	CA	PHE	79	-37.243	85.413	-26.181	1.00	45.00
	ATOM	639	CB	PHE	79	-36.660	85.542	-27.603	1.00	40.51
	ATOM	640	CG	PHE	79	-35.807	86.747	-27.816	1.00	39.52
50	ATOM	641	CD1	PHE	79	-34.424	86.623	-27.887	1.00	36.80
	ATOM	642	CD2	PHE	79	-36.382	88.003	-27.951	1.00	37.89
	ATOM	643	CE1	PHE	79	-33.624	87.735	-28.091	1.00	37.52
	ATOM	644	CE2	PHE	79	-35.588	89.120	-28.157	1.00	35.59
	ATOM	645	CZ	PHE	79	-34.206	88.987	-28.227	1.00	37.58
55	ATOM	646	C	PHE	79	-38.083	84.136	-26.205	1.00	43.50
	ATOM	647	O	PHE	79	-37.555	83.029	-26.325	1.00	37.85
	ATOM	648	N	GLY	80	-39.398	84.312	-26.131	1.00	46.65
	ATOM	649	CA	GLY	80	-40.321	83.195	-26.182	1.00	48.38
	ATOM	650	C	GLY	80	-40.009	81.989	-25.324	1.00	47.89
60	ATOM	651	O	GLY	80	-39.715	82.116	-24.132	1.00	51.15
	ATOM	652	N	ASP	81	-40.073	80.810	-25.939	1.00	45.07
	ATOM	653	CA	ASP	81	-39.832	79.567	-25.230	1.00	43.99
	ATOM	654	CB	ASP	81	-40.830	78.496	-25.691	1.00	45.66
	ATOM	655	CG	ASP	81	-40.587	78.036	-27.120	1.00	47.49
65	ATOM	656	OD1	ASP	81	-39.833	78.705	-27.861	1.00	51.90
	ATOM	657	OD2	ASP	81	-41.163	77.002	-27.511	1.00	48.08

	ATOM	658	C	ASP	81	-38.412	79.043	-25.356	1.00	42.29
	ATOM	659	O	ASP	81	-38.176	77.846	-25.190	1.00	39.94
	ATOM	660	N	GLU	82	-37.468	79.931	-25.649	1.00	39.91
	ATOM	661	CA	GLU	82	-36.067	79.531	-25.756	1.00	36.58
5	ATOM	662	CB	GLU	82	-35.170	80.703	-26.157	1.00	34.91
	ATOM	663	CG	GLU	82	-35.145	81.090	-27.601	1.00	37.82
	ATOM	664	CD	GLU	82	-33.850	81.798	-27.961	1.00	40.75
	ATOM	665	OE1	GLU	82	-33.401	82.662	-27.181	1.00	38.15
	ATOM	666	OE2	GLU	82	-33.270	81.491	-29.023	1.00	47.99
10	ATOM	667	C	GLU	82	-35.573	79.066	-24.393	1.00	36.58
	ATOM	668	O	GLU	82	-36.030	79.552	-23.362	1.00	38.40
	ATOM	669	N	LEU	83	-34.640	78.125	-24.393	1.00	33.83
	ATOM	670	CA	LEU	83	-34.033	77.659	-23.157	1.00	32.70
	ATOM	671	CB	LEU	83	-33.528	76.224	-23.306	1.00	30.78
15	ATOM	672	CG	LEU	83	-34.462	75.089	-22.893	1.00	29.26
	ATOM	673	CD1	LEU	83	-35.867	75.344	-23.377	1.00	29.60
	ATOM	674	CD2	LEU	83	-33.932	73.793	-23.450	1.00	28.65
	ATOM	675	C	LEU	83	-32.860	78.605	-23.021	1.00	33.88
	ATOM	676	O	LEU	83	-32.143	78.832	-23.989	1.00	38.46
20	ATOM	677	N	SER	84	-32.663	79.174	-21.840	1.00	34.94
	ATOM	678	CA	SER	84	-31.564	80.111	-21.659	1.00	39.77
	ATOM	679	CB	SER	84	-31.855	81.054	-20.493	1.00	38.92
	ATOM	680	OG	SER	84	-32.051	80.325	-19.306	1.00	43.11
	ATOM	681	C	SER	84	-30.216	79.428	-21.453	1.00	39.89
25	ATOM	682	O	SER	84	-29.167	80.070	-21.493	1.00	42.52
	ATOM	683	N	LEU	85	-30.247	78.121	-21.240	1.00	40.92
	ATOM	684	CA	LEU	85	-29.029	77.357	-21.045	1.00	34.66
	ATOM	685	CB	LEU	85	-29.132	76.529	-19.764	1.00	35.12
	ATOM	686	CG	LEU	85	-27.906	75.834	-19.155	1.00	36.92
30	ATOM	687	CD1	LEU	85	-27.442	74.711	-20.050	1.00	38.24
	ATOM	688	CD2	LEU	85	-26.795	76.850	-18.936	1.00	36.08
	ATOM	689	C	LEU	85	-28.871	76.453	-22.253	1.00	35.30
	ATOM	690	O	LEU	85	-29.689	75.572	-22.501	1.00	32.85
	ATOM	691	N	VAL	86	-27.821	76.695	-23.023	1.00	33.26
35	ATOM	692	CA	VAL	86	-27.550	75.900	-24.210	1.00	34.57
	ATOM	693	CB	VAL	86	-27.401	76.787	-25.480	1.00	35.39
	ATOM	694	CG1	VAL	86	-26.950	75.946	-26.659	1.00	31.44
	ATOM	695	CG2	VAL	86	-28.719	77.462	-25.802	1.00	38.08
	ATOM	696	C	VAL	86	-26.252	75.153	-24.017	1.00	32.85
40	ATOM	697	O	VAL	86	-25.260	75.728	-23.576	1.00	35.10
	ATOM	698	N	THR	87	-26.246	73.869	-24.328	1.00	33.41
	ATOM	699	CA	THR	87	-25.011	73.133	-24.206	1.00	33.56
	ATOM	700	CB	THR	87	-25.219	71.775	-23.467	1.00	32.70
	ATOM	701	OG1	THR	87	-24.444	70.746	-24.095	1.00	34.29
45	ATOM	702	CG2	THR	87	-26.678	71.404	-23.427	1.00	34.95
	ATOM	703	C	THR	87	-24.371	72.948	-25.587	1.00	30.57
	ATOM	704	O	THR	87	-24.936	72.317	-26.477	1.00	28.08
	ATOM	705	N	LEU	88	-23.212	73.577	-25.759	1.00	28.42
	ATOM	706	CA	LEU	88	-22.435	73.486	-26.981	1.00	31.31
50	ATOM	707	CB	LEU	88	-21.618	74.765	-27.214	1.00	28.21
	ATOM	708	CG	LEU	88	-22.079	76.226	-27.244	1.00	29.12
	ATOM	709	CD1	LEU	88	-23.562	76.332	-27.207	1.00	31.12
	ATOM	710	CD2	LEU	88	-21.457	76.962	-26.082	1.00	28.72
	ATOM	711	C	LEU	88	-21.430	72.346	-26.750	1.00	41.15
55	ATOM	712	O	LEU	88	-20.988	72.105	-25.624	1.00	52.30
	ATOM	713	N	PHE	89	-21.071	71.616	-27.790	1.00	39.23
	ATOM	714	CA	PHE	89	-20.048	70.583	-27.618	1.00	42.27
	ATOM	715	CB	PHE	89	-18.744	71.289	-27.235	1.00	35.63
	ATOM	716	CG	PHE	89	-18.523	72.554	-28.022	1.00	35.06
60	ATOM	717	CD1	PHE	89	-18.080	73.715	-27.395	1.00	31.56
	ATOM	718	CD2	PHE	89	-18.884	72.614	-29.379	1.00	30.96
	ATOM	719	CE1	PHE	89	-18.014	74.920	-28.102	1.00	33.76
	ATOM	720	CE2	PHE	89	-18.824	73.807	-30.090	1.00	27.38
	ATOM	721	CZ	PHE	89	-18.393	74.963	-29.452	1.00	30.29
65	ATOM	722	C	PHE	89	-20.340	69.376	-26.698	1.00	39.99
	ATOM	723	O	PHE	89	-21.250	68.592	-26.985	1.00	45.34

	ATOM	724	N	ARG	90	-19.589	69.184	-25.624	1.00	38.92
	ATOM	725	CA	ARG	90	-19.826	67.997	-24.782	1.00	41.66
	ATOM	726	CB	ARG	90	-21.336	67.749	-24.589	1.00	39.24
	ATOM	727	CG	ARG	90	-21.753	66.265	-24.560	1.00	29.41
5	ATOM	728	CD	ARG	90	-23.267	66.085	-24.337	1.00	31.13
	ATOM	729	NE	ARG	90	-23.706	64.709	-24.582	1.00	31.75
	ATOM	730	CZ	ARG	90	-24.514	64.328	-25.575	1.00	28.86
	ATOM	731	NH1	ARG	90	-25.002	65.208	-26.434	1.00	25.48
	ATOM	732	NH2	ARG	90	-24.814	63.049	-25.729	1.00	29.84
10	ATOM	733	C	ARG	90	-19.204	66.696	-25.326	1.00	39.23
	ATOM	734	O	ARG	90	-19.400	66.343	-26.488	1.00	31.33
	ATOM	735	N	CYS	91	-18.453	65.989	-24.478	1.00	40.12
	ATOM	736	CA	CYS	91	-17.865	64.722	-24.883	1.00	37.09
	ATOM	737	C	CYS	91	-17.959	63.593	-23.896	1.00	37.21
15	ATOM	738	O	CYS	91	-18.286	63.793	-22.733	1.00	37.46
	ATOM	739	CB	CYS	91	-16.419	64.867	-25.299	1.00	40.13
	ATOM	740	SG	CYS	91	-15.117	65.454	-24.158	1.00	41.64
	ATOM	741	N	ILE	92	-17.660	62.395	-24.387	1.00	34.02
	ATOM	742	CA	ILE	92	-17.741	61.185	-23.582	1.00	33.22
20	ATOM	743	CB	ILE	92	-19.008	60.387	-23.951	1.00	32.88
	ATOM	744	CG2	ILE	92	-19.199	59.235	-22.989	1.00	32.61
	ATOM	745	CG1	ILE	92	-20.229	61.309	-23.904	1.00	34.64
	ATOM	746	CD1	ILE	92	-21.509	60.686	-24.437	1.00	32.30
	ATOM	747	C	ILE	92	-16.534	60.271	-23.760	1.00	30.78
25	ATOM	748	O	ILE	92	-15.889	60.288	-24.788	1.00	33.43
	ATOM	749	N	GLN	93	-16.224	59.491	-22.732	1.00	29.35
	ATOM	750	CA	GLN	93	-15.125	58.540	-22.777	1.00	28.69
	ATOM	751	CB	GLN	93	-13.859	59.098	-22.126	1.00	28.17
	ATOM	752	CG	GLN	93	-12.953	59.976	-22.994	1.00	25.87
30	ATOM	753	CD	GLN	93	-12.403	59.271	-24.214	1.00	24.22
	ATOM	754	OE1	GLN	93	-13.022	59.272	-25.267	1.00	28.90
	ATOM	755	NE2	GLN	93	-11.234	58.661	-24.076	1.00	25.29
	ATOM	756	C	GLN	93	-15.555	57.324	-21.988	1.00	31.44
	ATOM	757	O	GLN	93	-15.971	57.451	-20.846	1.00	32.07
35	ATOM	758	N	ASN	94	-15.491	56.147	-22.595	1.00	35.77
	ATOM	759	CA	ASN	94	-15.835	54.940	-21.869	1.00	33.56
	ATOM	760	CB	ASN	94	-15.932	53.740	-22.813	1.00	34.70
	ATOM	761	CG	ASN	94	-17.260	53.664	-23.524	1.00	33.01
	ATOM	762	OD1	ASN	94	-18.306	53.712	-22.899	1.00	30.56
40	ATOM	763	ND2	ASN	94	-17.223	53.526	-24.834	1.00	36.61
	ATOM	764	C	ASN	94	-14.692	54.737	-20.874	1.00	35.80
	ATOM	765	O	ASN	94	-13.532	55.016	-21.182	1.00	36.63
	ATOM	766	N	MET	95	-15.015	54.264	-19.677	1.00	35.73
	ATOM	767	CA	MET	95	-13.996	54.051	-18.657	1.00	34.73
45	ATOM	768	CB	MET	95	-14.404	54.766	-17.366	1.00	32.97
	ATOM	769	CG	MET	95	-14.718	56.247	-17.521	1.00	27.35
	ATOM	770	SD	MET	95	-13.391	57.202	-18.283	1.00	25.14
	ATOM	771	CE	MET	95	-12.179	57.271	-17.008	1.00	15.55
	ATOM	772	C	MET	95	-13.756	52.568	-18.368	1.00	36.36
50	ATOM	773	O	MET	95	-14.653	51.739	-18.548	1.00	40.70
	ATOM	774	N	PRO	96	-12.531	52.212	-17.944	1.00	36.06
	ATOM	775	CD	PRO	96	-11.322	53.027	-18.065	1.00	35.32
	ATOM	776	CA	PRO	96	-12.168	50.833	-17.617	1.00	40.00
	ATOM	777	CB	PRO	96	-10.655	50.805	-17.810	1.00	33.97
55	ATOM	778	CG	PRO	96	-10.360	52.020	-18.594	1.00	37.05
	ATOM	779	C	PRO	96	-12.531	50.611	-16.151	1.00	46.34
	ATOM	780	O	PRO	96	-12.814	51.567	-15.431	1.00	47.59
	ATOM	781	N	GLU	97	-12.509	49.361	-15.698	1.00	52.14
	ATOM	782	CA	GLU	97	-12.843	49.063	-14.311	1.00	57.27
60	ATOM	783	CB	GLU	97	-13.369	47.633	-14.184	1.00	62.29
	ATOM	784	CG	GLU	97	-14.246	47.387	-12.960	1.00	72.31
	ATOM	785	CD	GLU	97	-15.742	47.418	-13.301	1.00	79.34
	ATOM	786	OE1	GLU	97	-16.195	46.557	-14.118	1.00	84.94
	ATOM	787	OE2	GLU	97	-16.462	48.303	-12.759	1.00	81.39
65	ATOM	788	C	GLU	97	-11.591	49.212	-13.454	1.00	56.67
	ATOM	789	O	GLU	97	-11.656	49.667	-12.308	1.00	59.18

	ATOM	790	N	THR	98	-10.450	48.845	-14.025	1.00	57.55
	ATOM	791	CA	THR	98	-9.177	48.888	-13.315	1.00	59.71
	ATOM	792	CB	THR	98	-8.099	48.162	-14.110	1.00	59.24
	ATOM	793	OG1	THR	98	-7.980	48.792	-15.400	1.00	68.81
5	ATOM	794	CG2	THR	98	-8.464	46.671	-14.261	1.00	56.15
	ATOM	795	C	THR	98	-8.629	50.266	-12.946	1.00	60.32
	ATOM	796	O	THR	98	-9.002	50.822	-11.906	1.00	65.03
	ATOM	797	N	LEU	99	-7.735	50.806	-13.776	1.00	54.94
	ATOM	798	CA	LEU	99	-7.122	52.108	-13.500	1.00	51.13
10	ATOM	799	CB	LEU	99	-5.605	51.997	-13.635	1.00	50.56
	ATOM	800	CG	LEU	99	-4.873	51.250	-12.524	1.00	52.80
	ATOM	801	CD1	LEU	99	-3.476	50.844	-12.989	1.00	48.38
	ATOM	802	CD2	LEU	99	-4.804	52.138	-11.291	1.00	50.80
	ATOM	803	C	LEU	99	-7.633	53.241	-14.395	1.00	50.20
15	ATOM	804	O	LEU	99	-6.950	53.661	-15.340	1.00	50.50
	ATOM	805	N	PRO	100	-8.827	53.778	-14.086	1.00	47.97
	ATOM	806	CD	PRO	100	-9.638	53.512	-12.888	1.00	45.68
	ATOM	807	CA	PRO	100	-9.422	54.861	-14.875	1.00	46.54
	ATOM	808	CB	PRO	100	-10.666	55.239	-14.070	1.00	45.50
20	ATOM	809	CG	PRO	100	-10.994	53.985	-13.327	1.00	45.80
	ATOM	810	C	PRO	100	-8.487	56.045	-15.057	1.00	46.78
	ATOM	811	O	PRO	100	-8.014	56.627	-14.088	1.00	52.13
	ATOM	812	N	ASN	101	-8.224	56.396	-16.304	1.00	42.71
	ATOM	813	CA	ASN	101	-7.373	57.525	-16.623	1.00	41.37
25	ATOM	814	CB	ASN	101	-5.910	57.176	-16.379	1.00	43.24
	ATOM	815	CG	ASN	101	-5.498	57.362	-14.933	1.00	45.93
	ATOM	816	OD1	ASN	101	-5.410	58.488	-14.438	1.00	45.33
	ATOM	817	ND2	ASN	101	-5.248	56.252	-14.239	1.00	48.11
	ATOM	818	C	ASN	101	-7.588	57.879	-18.081	1.00	42.66
30	ATOM	819	O	ASN	101	-6.817	57.469	-18.951	1.00	44.79
	ATOM	820	N	ASN	102	-8.644	58.635	-18.365	1.00	40.46
	ATOM	821	CA	ASN	102	-8.892	58.991	-19.743	1.00	37.90
	ATOM	822	CB	ASN	102	-10.256	58.487	-20.187	1.00	34.01
	ATOM	823	CG	ASN	102	-10.197	57.061	-20.697	1.00	32.44
35	ATOM	824	OD1	ASN	102	-9.259	56.685	-21.395	1.00	29.07
	ATOM	825	ND2	ASN	102	-11.198	56.263	-20.358	1.00	33.79
	ATOM	826	C	ASN	102	-8.705	60.431	-20.180	1.00	39.20
	ATOM	827	O	ASN	102	-7.831	60.707	-20.993	1.00	46.07
	ATOM	828	N	SER	103	-9.484	61.363	-19.669	1.00	35.42
40	ATOM	829	CA	SER	103	-9.318	62.752	-20.134	1.00	40.28
	ATOM	830	CB	SER	103	-7.850	63.229	-20.007	1.00	40.68
	ATOM	831	OG	SER	103	-7.157	63.221	-21.251	1.00	31.76
	ATOM	832	C	SER	103	-9.791	62.896	-21.600	1.00	35.24
	ATOM	833	O	SER	103	-9.315	62.216	-22.508	1.00	24.65
45	ATOM	834	N	CYS	104	-10.762	63.776	-21.798	1.00	32.88
	ATOM	835	CA	CYS	104	-11.317	64.023	-23.104	1.00	31.77
	ATOM	836	C	CYS	104	-11.232	65.534	-23.381	1.00	30.58
	ATOM	837	O	CYS	104	-11.529	66.353	-22.516	1.00	29.23
	ATOM	838	CB	CYS	104	-12.786	63.521	-23.176	1.00	28.43
50	ATOM	839	SG	CYS	104	-13.533	64.197	-24.669	1.00	46.07
	ATOM	840	N	TYR	105	-10.790	65.896	-24.584	1.00	30.34
	ATOM	841	CA	TYR	105	-10.695	67.294	-25.002	1.00	26.12
	ATOM	842	CB	TYR	105	-9.315	67.588	-25.588	1.00	27.03
	ATOM	843	CG	TYR	105	-9.171	68.959	-26.229	1.00	25.61
55	ATOM	844	CD1	TYR	105	-8.605	70.032	-25.533	1.00	24.90
	ATOM	845	CE1	TYR	105	-8.475	71.283	-26.117	1.00	26.08
	ATOM	846	CD2	TYR	105	-9.605	69.188	-27.531	1.00	25.41
	ATOM	847	CE2	TYR	105	-9.482	70.439	-28.120	1.00	28.91
	ATOM	848	CZ	TYR	105	-8.916	71.481	-27.409	1.00	27.58
60	ATOM	849	OH	TYR	105	-8.783	72.717	-28.002	1.00	29.89
	ATOM	850	C	TYR	105	-11.751	67.543	-26.075	1.00	29.20
	ATOM	851	O	TYR	105	-12.032	66.675	-26.895	1.00	33.10
	ATOM	852	N	SER	106	-12.340	68.728	-26.074	1.00	30.48
	ATOM	853	CA	SER	106	-13.348	69.049	-27.072	1.00	29.22
65	ATOM	854	CB	SER	106	-14.709	68.496	-26.639	1.00	28.30
	ATOM	855	OG	SER	106	-15.706	68.745	-27.610	1.00	28.62

	ATOM	856	C	SER	106	-13.407	70.560	-27.210	1.00	29.94
	ATOM	857	O	SER	106	-13.255	71.271	-26.225	1.00	29.72
	ATOM	858	N	ALA	107	-13.603	71.047	-28.433	1.00	27.64
	ATOM	859	CA	ALA	107	-13.679	72.481	-28.679	1.00	24.33
5	ATOM	860	CB	ALA	107	-12.284	73.059	-28.838	1.00	21.30
	ATOM	861	C	ALA	107	-14.514	72.800	-29.909	1.00	25.56
	ATOM	862	O	ALA	107	-14.816	71.930	-30.715	1.00	28.03
	ATOM	863	N	GLY	108	-14.885	74.063	-30.047	1.00	26.07
	ATOM	864	CA	GLY	108	-15.671	74.482	-31.188	1.00	25.49
10	ATOM	865	C	GLY	108	-15.881	75.978	-31.159	1.00	29.03
	ATOM	866	O	GLY	108	-15.471	76.638	-30.214	1.00	32.20
	ATOM	867	N	ILE	109	-16.520	76.517	-32.188	1.00	29.34
	ATOM	868	CA	ILE	109	-16.778	77.947	-32.262	1.00	27.71
	ATOM	869	CB	ILE	109	-16.390	78.503	-33.642	1.00	27.48
15	ATOM	870	CG2	ILE	109	-16.684	79.987	-33.712	1.00	26.46
	ATOM	871	CG1	ILE	109	-14.907	78.242	-33.897	1.00	26.16
	ATOM	872	CD1	ILE	109	-14.453	78.594	-35.285	1.00	29.07
	ATOM	873	C	ILE	109	-18.249	78.216	-32.015	1.00	28.66
	ATOM	874	O	ILE	109	-19.109	77.448	-32.440	1.00	30.59
20	ATOM	875	N	ALA	110	-18.543	79.305	-31.318	1.00	25.60
	ATOM	876	CA	ALA	110	-19.922	79.674	-31.022	1.00	26.36
	ATOM	877	CB	ALA	110	-20.347	79.072	-29.703	1.00	20.26
	ATOM	878	C	ALA	110	-20.051	81.185	-30.957	1.00	29.61
	ATOM	879	O	ALA	110	-19.083	81.882	-30.689	1.00	27.61
25	ATOM	880	N	LYS	111	-21.242	81.702	-31.223	1.00	33.77
	ATOM	881	CA	LYS	111	-21.432	83.132	-31.141	1.00	35.09
	ATOM	882	CB	LYS	111	-22.382	83.635	-32.220	1.00	39.38
	ATOM	883	CG	LYS	111	-22.700	85.111	-32.020	1.00	49.74
	ATOM	884	CD	LYS	111	-22.888	85.858	-33.312	1.00	53.95
30	ATOM	885	CE	LYS	111	-23.027	87.348	-33.037	1.00	54.28
	ATOM	886	NZ	LYS	111	-23.243	88.109	-34.314	1.00	60.85
	ATOM	887	C	LYS	111	-21.988	83.455	-29.759	1.00	35.28
	ATOM	888	O	LYS	111	-22.989	82.889	-29.337	1.00	36.58
	ATOM	889	N	LEU	112	-21.319	84.362	-29.061	1.00	32.32
35	ATOM	890	CA	LEU	112	-21.720	84.755	-27.724	1.00	29.67
	ATOM	891	CB	LEU	112	-20.605	84.428	-26.733	1.00	27.04
	ATOM	892	CG	LEU	112	-20.033	83.014	-26.802	1.00	24.38
	ATOM	893	CD1	LEU	112	-18.852	82.891	-25.860	1.00	27.56
	ATOM	894	CD2	LEU	112	-21.101	82.013	-26.454	1.00	19.52
40	ATOM	895	C	LEU	112	-22.016	86.246	-27.681	1.00	32.58
	ATOM	896	O	LEU	112	-21.649	86.986	-28.593	1.00	31.51
	ATOM	897	N	GLU	113	-22.674	86.686	-26.615	1.00	35.75
	ATOM	898	CA	GLU	113	-23.016	88.088	-26.461	1.00	40.08
	ATOM	899	CB	GLU	113	-24.506	88.306	-26.650	1.00	44.23
45	ATOM	900	CG	GLU	113	-25.116	87.642	-27.849	1.00	50.12
	ATOM	901	CD	GLU	113	-26.414	88.317	-28.228	1.00	58.08
	ATOM	902	OE1	GLU	113	-27.176	88.717	-27.299	1.00	61.13
	ATOM	903	OE2	GLU	113	-26.668	88.449	-29.454	1.00	63.64
	ATOM	904	C	GLU	113	-22.667	88.608	-25.089	1.00	41.63
50	ATOM	905	O	GLU	113	-22.507	87.840	-24.149	1.00	48.29
	ATOM	906	N	GLU	114	-22.565	89.929	-24.987	1.00	40.97
	ATOM	907	CA	GLU	114	-22.279	90.601	-23.726	1.00	36.68
	ATOM	908	CB	GLU	114	-22.582	92.080	-23.842	1.00	41.19
	ATOM	909	CG	GLU	114	-21.435	92.974	-24.147	1.00	46.58
55	ATOM	910	CD	GLU	114	-21.701	94.367	-23.619	1.00	51.12
	ATOM	911	OE1	GLU	114	-21.761	94.520	-22.375	1.00	51.88
	ATOM	912	OE2	GLU	114	-21.864	95.299	-24.444	1.00	57.08
	ATOM	913	C	GLU	114	-23.212	90.058	-22.664	1.00	35.51
	ATOM	914	O	GLU	114	-24.426	90.125	-22.819	1.00	38.92
60	ATOM	915	N	GLY	115	-22.658	89.538	-21.580	1.00	32.99
	ATOM	916	CA	GLY	115	-23.514	89.035	-20.529	1.00	31.79
	ATOM	917	C	GLY	115	-23.658	87.537	-20.519	1.00	34.39
	ATOM	918	O	GLY	115	-24.093	86.975	-19.523	1.00	39.20
	ATOM	919	N	ASP	116	-23.322	86.888	-21.626	1.00	35.66
65	ATOM	920	CA	ASP	116	-23.403	85.438	-21.682	1.00	38.09
	ATOM	921	CB	ASP	116	-23.131	84.919	-23.099	1.00	36.00

	ATOM	922	CG	ASP	116	-24.310	85.093	-24.030	1.00	35.09
	ATOM	923	OD1	ASP	116	-25.436	85.340	-23.550	1.00	33.81
	ATOM	924	OD2	ASP	116	-24.112	84.964	-25.252	1.00	35.24
	ATOM	925	C	ASP	116	-22.338	84.887	-20.745	1.00	39.38
5	ATOM	926	O	ASP	116	-21.321	85.540	-20.493	1.00	37.91
	ATOM	927	N	GLU	117	-22.574	83.693	-20.217	1.00	41.54
	ATOM	928	CA	GLU	117	-21.597	83.062	-19.341	1.00	37.92
	ATOM	929	CB	GLU	117	-22.124	82.967	-17.914	1.00	38.90
	ATOM	930	CG	GLU	117	-22.493	84.296	-17.306	1.00	48.15
10	ATOM	931	CD	GLU	117	-22.971	84.156	-15.876	1.00	52.62
	ATOM	932	OE1	GLU	117	-23.676	83.155	-15.571	1.00	55.49
	ATOM	933	OE2	GLU	117	-22.646	85.055	-15.062	1.00	57.10
	ATOM	934	C	GLU	117	-21.326	81.667	-19.869	1.00	34.02
	ATOM	935	O	GLU	117	-22.226	81.014	-20.395	1.00	33.31
15	ATOM	936	N	LEU	118	-20.081	81.225	-19.756	1.00	30.68
	ATOM	937	CA	LEU	118	-19.700	79.893	-20.194	1.00	27.14
	ATOM	938	CB	LEU	118	-18.480	79.957	-21.113	1.00	26.49
	ATOM	939	CG	LEU	118	-18.626	80.646	-22.465	1.00	26.21
	ATOM	940	CD1	LEU	118	-17.285	80.668	-23.162	1.00	23.05
20	ATOM	941	CD2	LEU	118	-19.652	79.919	-23.309	1.00	25.38
	ATOM	942	C	LEU	118	-19.347	79.069	-18.963	1.00	28.33
	ATOM	943	O	LEU	118	-18.805	79.599	-17.991	1.00	31.46
	ATOM	944	N	GLN	119	-19.661	77.778	-18.989	1.00	28.96
	ATOM	945	CA	GLN	119	-19.319	76.910	-17.869	1.00	29.13
25	ATOM	946	CB	GLN	119	-20.429	76.928	-16.828	1.00	28.23
	ATOM	947	CG	GLN	119	-21.690	76.247	-17.266	1.00	36.38
	ATOM	948	CD	GLN	119	-22.783	76.348	-16.230	1.00	36.57
	ATOM	949	OE1	GLN	119	-23.771	75.617	-16.291	1.00	40.70
	ATOM	950	NE2	GLN	119	-22.622	77.260	-15.279	1.00	36.80
30	ATOM	951	C	GLN	119	-19.038	75.480	-18.319	1.00	29.43
	ATOM	952	O	GLN	119	-19.535	75.033	-19.350	1.00	28.80
	ATOM	953	N	LEU	120	-18.219	74.784	-17.538	1.00	32.26
	ATOM	954	CA	LEU	120	-17.828	73.402	-17.801	1.00	29.96
	ATOM	955	CB	LEU	120	-16.303	73.270	-17.675	1.00	31.17
35	ATOM	956	CG	LEU	120	-15.482	72.057	-18.131	1.00	30.26
	ATOM	957	CD1	LEU	120	-16.105	70.767	-17.661	1.00	27.10
	ATOM	958	CD2	LEU	120	-15.370	72.085	-19.628	1.00	30.96
	ATOM	959	C	LEU	120	-18.517	72.542	-16.739	1.00	30.51
	ATOM	960	O	LEU	120	-18.211	72.640	-15.550	1.00	32.88
40	ATOM	961	N	ALA	121	-19.442	71.694	-17.173	1.00	27.40
	ATOM	962	CA	ALA	121	-20.180	70.840	-16.253	1.00	25.55
	ATOM	963	CB	ALA	121	-21.655	71.208	-16.298	1.00	19.41
	ATOM	964	C	ALA	121	-20.016	69.342	-16.511	1.00	29.89
	ATOM	965	O	ALA	121	-20.014	68.892	-17.656	1.00	35.88
45	ATOM	966	N	ILE	122	-19.872	68.579	-15.432	1.00	30.33
	ATOM	967	CA	ILE	122	-19.743	67.130	-15.523	1.00	30.27
	ATOM	968	CB	ILE	122	-18.583	66.612	-14.668	1.00	29.27
	ATOM	969	CG2	ILE	122	-18.437	65.121	-14.870	1.00	30.38
	ATOM	970	CG1	ILE	122	-17.287	67.328	-15.059	1.00	26.03
50	ATOM	971	CD1	ILE	122	-16.068	66.918	-14.253	1.00	29.21
	ATOM	972	C	ILE	122	-21.055	66.562	-14.990	1.00	33.20
	ATOM	973	O	ILE	122	-21.372	66.722	-13.813	1.00	33.30
	ATOM	974	N	PRO	123	-21.837	65.895	-15.856	1.00	34.70
	ATOM	975	CD	PRO	123	-21.551	65.696	-17.280	1.00	34.40
55	ATOM	976	CA	PRO	123	-23.133	65.294	-15.512	1.00	36.49
	ATOM	977	CB	PRO	123	-23.749	64.974	-16.878	1.00	30.22
	ATOM	978	CG	PRO	123	-22.926	65.758	-17.858	1.00	36.72
	ATOM	979	C	PRO	123	-23.033	64.036	-14.652	1.00	39.64
	ATOM	980	O	PRO	123	-23.538	62.978	-15.040	1.00	40.48
60	ATOM	981	N	ARG	124	-22.390	64.150	-13.493	1.00	44.58
	ATOM	982	CA	ARG	124	-22.233	63.018	-12.591	1.00	48.04
	ATOM	983	CB	ARG	124	-20.963	62.245	-12.927	1.00	52.91
	ATOM	984	CG	ARG	124	-20.884	61.795	-14.374	1.00	60.50
	ATOM	985	CD	ARG	124	-21.068	60.295	-14.433	1.00	72.57
65	ATOM	986	NE	ARG	124	-21.167	59.804	-15.819	1.00	84.90
	ATOM	987	CZ	ARG	124	-22.306	59.657	-16.508	1.00	85.44

	ATOM	988	NH1	ARG	124	-23.475	59.966	-15.946	1.00	87.69
	ATOM	989	NH2	ARG	124	-22.272	59.202	-17.768	1.00	82.78
	ATOM	990	C	ARG	124	-22.199	63.457	-11.136	1.00	48.39
	ATOM	991	O	ARG	124	-21.715	64.544	-10.820	1.00	45.26
5	ATOM	992	N	GLU	125	-22.710	62.589	-10.267	1.00	52.84
	ATOM	993	CA	GLU	125	-22.791	62.839	-8.830	1.00	55.02
	ATOM	994	CB	GLU	125	-23.264	61.570	-8.115	1.00	63.62
	ATOM	995	CG	GLU	125	-22.867	60.271	-8.821	1.00	70.62
	ATOM	996	CD	GLU	125	-23.332	60.228	-10.272	1.00	74.76
10	ATOM	997	OE1	GLU	125	-24.570	60.210	-10.506	1.00	75.75
	ATOM	998	OE2	GLU	125	-22.458	60.213	-11.174	1.00	80.71
	ATOM	999	C	GLU	125	-21.508	63.346	-8.185	1.00	53.44
	ATOM	1000	O	GLU	125	-21.441	64.505	-7.774	1.00	57.50
	ATOM	1001	N	ASN	126	-20.507	62.488	-8.047	1.00	47.18
15	ATOM	1002	CA	ASN	126	-19.248	62.937	-7.469	1.00	47.59
	ATOM	1003	CB	ASN	126	-19.042	62.388	-6.059	1.00	51.03
	ATOM	1004	CG	ASN	126	-19.352	63.422	-4.986	1.00	57.07
	ATOM	1005	OD1	ASN	126	-20.521	63.719	-4.696	1.00	61.15
	ATOM	1006	ND2	ASN	126	-18.299	63.994	-4.402	1.00	56.63
20	ATOM	1007	C	ASN	126	-18.139	62.476	-8.375	1.00	47.06
	ATOM	1008	O	ASN	126	-17.358	61.581	-8.042	1.00	45.61
	ATOM	1009	N	ALA	127	-18.095	63.105	-9.542	1.00	43.50
	ATOM	1010	CA	ALA	127	-17.121	62.781	-10.560	1.00	38.26
	ATOM	1011	CB	ALA	127	-17.147	63.841	-11.636	1.00	37.97
25	ATOM	1012	C	ALA	127	-15.720	62.645	-9.998	1.00	37.16
	ATOM	1013	O	ALA	127	-15.231	63.524	-9.290	1.00	33.14
	ATOM	1014	N	GLN	128	-15.085	61.519	-10.302	1.00	37.87
	ATOM	1015	CA	GLN	128	-13.711	61.292	-9.877	1.00	37.19
	ATOM	1016	CB	GLN	128	-13.456	59.796	-9.731	1.00	40.23
30	ATOM	1017	CG	GLN	128	-14.219	59.200	-8.571	1.00	38.14
	ATOM	1018	CD	GLN	128	-13.930	59.944	-7.288	1.00	39.51
	ATOM	1019	OE1	GLN	128	-12.812	59.895	-6.763	1.00	40.36
	ATOM	1020	NE2	GLN	128	-14.931	60.664	-6.783	1.00	41.53
	ATOM	1021	C	GLN	128	-12.861	61.909	-10.984	1.00	35.55
35	ATOM	1022	O	GLN	128	-12.652	61.323	-12.050	1.00	35.57
	ATOM	1023	N	ILE	129	-12.387	63.114	-10.705	1.00	34.64
	ATOM	1024	CA	ILE	129	-11.629	63.919	-11.653	1.00	29.60
	ATOM	1025	CB	ILE	129	-12.491	65.191	-11.983	1.00	31.03
	ATOM	1026	CG2	ILE	129	-11.686	66.463	-11.916	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-13.166	65.004	-13.324	1.00	26.37
	ATOM	1028	CD1	ILE	129	-14.081	63.838	-13.346	1.00	34.70
	ATOM	1029	C	ILE	129	-10.253	64.320	-11.130	1.00	29.55
	ATOM	1030	O	ILE	129	-10.037	64.386	-9.929	1.00	33.42
	ATOM	1031	N	SER	130	-9.316	64.577	-12.036	1.00	27.67
45	ATOM	1032	CA	SER	130	-7.987	65.024	-11.629	1.00	25.00
	ATOM	1033	CB	SER	130	-6.922	64.533	-12.594	1.00	22.09
	ATOM	1034	OG	SER	130	-5.684	65.172	-12.315	1.00	20.89
	ATOM	1035	C	SER	130	-7.989	66.544	-11.641	1.00	26.03
	ATOM	1036	O	SER	130	-8.500	67.145	-12.570	1.00	30.77
50	ATOM	1037	N	LEU	131	-7.416	67.169	-10.624	1.00	24.49
	ATOM	1038	CA	LEU	131	-7.397	68.623	-10.562	1.00	28.07
	ATOM	1039	CB	LEU	131	-7.912	69.102	-9.205	1.00	28.87
	ATOM	1040	CG	LEU	131	-9.421	69.312	-9.084	1.00	30.76
	ATOM	1041	CD1	LEU	131	-10.177	68.140	-9.637	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-9.765	69.526	-7.633	1.00	33.37
	ATOM	1043	C	LEU	131	-6.044	69.254	-10.843	1.00	33.66
	ATOM	1044	O	LEU	131	-5.730	70.319	-10.312	1.00	37.67
	ATOM	1045	N	ASP	132	-5.242	68.598	-11.676	1.00	34.97
	ATOM	1046	CA	ASP	132	-3.934	69.130	-12.043	1.00	34.31
60	ATOM	1047	CB	ASP	132	-2.997	68.011	-12.494	1.00	42.68
	ATOM	1048	CG	ASP	132	-2.453	67.197	-11.331	1.00	47.98
	ATOM	1049	OD1	ASP	132	-1.845	66.128	-11.585	1.00	50.41
	ATOM	1050	OD2	ASP	132	-2.627	67.637	-10.167	1.00	46.42
	ATOM	1051	C	ASP	132	-4.125	70.134	-13.167	1.00	34.32
65	ATOM	1052	O	ASP	132	-4.833	69.871	-14.133	1.00	32.87
	ATOM	1053	N	GLY	133	-3.487	71.288	-13.026	1.00	34.53

	ATOM	1054	CA	GLY	133	-3.615	72.338	-14.017	1.00	36.93
	ATOM	1055	C	GLY	133	-3.305	71.978	-15.456	1.00	35.06
	ATOM	1056	O	GLY	133	-3.765	72.654	-16.376	1.00	40.44
	ATOM	1057	N	ASP	134	-2.536	70.920	-15.670	1.00	31.08
5	ATOM	1058	CA	ASP	134	-2.186	70.547	-17.026	1.00	27.63
	ATOM	1059	CB	ASP	134	-0.740	70.049	-17.076	1.00	28.43
	ATOM	1060	CG	ASP	134	-0.499	68.836	-16.205	1.00	31.78
	ATOM	1061	OD1	ASP	134	-1.133	68.728	-15.139	1.00	35.68
	ATOM	1062	OD2	ASP	134	0.344	67.994	-16.578	1.00	32.53
10	ATOM	1063	C	ASP	134	-3.133	69.546	-17.663	1.00	25.08
	ATOM	1064	O	ASP	134	-3.204	69.458	-18.868	1.00	19.71
	ATOM	1065	N	VAL	135	-3.897	68.816	-16.866	1.00	26.84
	ATOM	1066	CA	VAL	135	-4.811	67.846	-17.451	1.00	23.72
	ATOM	1067	CB	VAL	135	-4.762	66.512	-16.703	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-3.444	65.842	-16.980	1.00	22.76
	ATOM	1069	CG2	VAL	135	-4.952	66.733	-15.218	1.00	22.12
	ATOM	1070	C	VAL	135	-6.260	68.306	-17.581	1.00	26.54
	ATOM	1071	O	VAL	135	-6.983	67.802	-18.430	1.00	29.26
	ATOM	1072	N	THR	136	-6.700	69.247	-16.748	1.00	26.39
20	ATOM	1073	CA	THR	136	-8.069	69.743	-16.885	1.00	28.40
	ATOM	1074	CB	THR	136	-9.026	69.141	-15.802	1.00	28.21
	ATOM	1075	OG1	THR	136	-9.279	70.102	-14.779	1.00	37.13
	ATOM	1076	CG2	THR	136	-8.426	67.902	-15.187	1.00	26.90
	ATOM	1077	C	THR	136	-8.090	71.279	-16.876	1.00	31.60
25	ATOM	1078	O	THR	136	-7.707	71.924	-15.896	1.00	27.29
	ATOM	1079	N	PHE	137	-8.525	71.850	-18.001	1.00	31.45
	ATOM	1080	CA	PHE	137	-8.565	73.297	-18.178	1.00	26.64
	ATOM	1081	CB	PHE	137	-7.261	73.744	-18.823	1.00	26.84
	ATOM	1082	CG	PHE	137	-6.803	72.853	-19.941	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-7.264	73.046	-21.239	1.00	27.77
	ATOM	1084	CD2	PHE	137	-5.903	71.828	-19.701	1.00	23.70
	ATOM	1085	CE1	PHE	137	-6.835	72.235	-22.277	1.00	27.32
	ATOM	1086	CE2	PHE	137	-5.470	71.012	-20.735	1.00	26.31
	ATOM	1087	CZ	PHE	137	-5.938	71.219	-22.024	1.00	29.43
35	ATOM	1088	C	PHE	137	-9.772	73.754	-18.994	1.00	28.06
	ATOM	1089	O	PHE	137	-10.422	72.945	-19.640	1.00	25.04
	ATOM	1090	N	PHE	138	-10.059	75.055	-18.964	1.00	29.81
	ATOM	1091	CA	PHE	138	-11.226	75.605	-19.649	1.00	29.93
	ATOM	1092	CB	PHE	138	-12.261	75.963	-18.583	1.00	27.47
40	ATOM	1093	CG	PHE	138	-13.618	76.268	-19.120	1.00	28.49
	ATOM	1094	CD1	PHE	138	-14.030	75.775	-20.353	1.00	27.65
	ATOM	1095	CD2	PHE	138	-14.499	77.052	-18.379	1.00	31.06
	ATOM	1096	CE1	PHE	138	-15.297	76.063	-20.841	1.00	30.90
	ATOM	1097	CE2	PHE	138	-15.770	77.348	-18.855	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-16.171	76.854	-20.088	1.00	27.41
	ATOM	1099	C	PHE	138	-10.909	76.798	-20.573	1.00	30.91
	ATOM	1100	O	PHE	138	-10.289	77.775	-20.169	1.00	24.94
	ATOM	1101	N	GLY	139	-11.393	76.678	-21.812	1.00	36.94
	ATOM	1102	CA	GLY	139	-11.164	77.608	-22.916	1.00	36.08
50	ATOM	1103	C	GLY	139	-11.524	79.066	-23.058	1.00	36.48
	ATOM	1104	O	GLY	139	-11.238	79.857	-22.178	1.00	43.61
	ATOM	1105	N	ALA	140	-12.086	79.414	-24.217	1.00	36.60
	ATOM	1106	CA	ALA	140	-12.511	80.779	-24.582	1.00	34.94
	ATOM	1107	CB	ALA	140	-13.203	81.446	-23.406	1.00	36.94
55	ATOM	1108	C	ALA	140	-11.456	81.740	-25.164	1.00	32.53
	ATOM	1109	O	ALA	140	-10.619	82.285	-24.452	1.00	27.96
	ATOM	1110	N	LEU	141	-11.538	81.956	-26.475	1.00	33.23
	ATOM	1111	CA	LEU	141	-10.638	82.851	-27.216	1.00	36.11
	ATOM	1112	CB	LEU	141	-9.563	82.029	-27.927	1.00	37.53
60	ATOM	1113	CG	LEU	141	-8.539	82.696	-28.851	1.00	38.87
	ATOM	1114	CD1	LEU	141	-7.467	81.689	-29.190	1.00	40.67
	ATOM	1115	CD2	LEU	141	-9.184	83.194	-30.129	1.00	37.53
	ATOM	1116	C	LEU	141	-11.468	83.621	-28.248	1.00	38.89
	ATOM	1117	O	LEU	141	-12.217	83.021	-29.016	1.00	39.25
65	ATOM	1118	N	LYS	142	-11.340	84.943	-28.285	1.00	42.86
	ATOM	1119	CA	LYS	142	-12.129	85.712	-29.242	1.00	42.96

	ATOM	1120	CB	LYS	142	-12.384	87.119	-28.711	1.00	41.80
	ATOM	1121	CG	LYS	142	-13.175	87.962	-29.683	1.00	46.88
	ATOM	1122	CD	LYS	142	-13.837	89.152	-29.022	1.00	49.58
	ATOM	1123	CE	LYS	142	-14.613	89.944	-30.062	1.00	51.07
5	ATOM	1124	NZ	LYS	142	-15.359	91.081	-29.469	1.00	56.20
	ATOM	1125	C	LYS	142	-11.539	85.797	-30.645	1.00	43.41
	ATOM	1126	O	LYS	142	-10.373	86.140	-30.821	1.00	44.98
	ATOM	1127	N	LEU	143	-12.357	85.485	-31.646	1.00	42.81
	ATOM	1128	CA	LEU	143	-11.925	85.529	-33.041	1.00	40.65
10	ATOM	1129	CB	LEU	143	-12.773	84.584	-33.890	1.00	36.56
	ATOM	1130	CG	LEU	143	-12.825	83.107	-33.502	1.00	36.17
	ATOM	1131	CD1	LEU	143	-13.769	82.379	-34.440	1.00	37.95
	ATOM	1132	CD2	LEU	143	-11.439	82.503	-33.568	1.00	27.38
	ATOM	1133	C	LEU	143	-12.068	86.940	-33.599	1.00	43.14
15	ATOM	1134	O	LEU	143	-12.961	87.691	-33.190	1.00	46.28
	ATOM	1135	N	LEU	144	-11.197	87.305	-34.531	1.00	42.63
	ATOM	1136	CA	LEU	144	-11.271	88.628	-35.133	1.00	44.11
	ATOM	1137	CB	LEU	144	-9.940	88.999	-35.779	1.00	44.88
	ATOM	1138	CG	LEU	144	-8.782	89.153	-34.796	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-7.504	89.418	-35.544	1.00	47.71
	ATOM	1140	CD2	LEU	144	-9.075	90.290	-33.833	1.00	47.87
	ATOM	1141	C	LEU	144	-12.361	88.653	-36.186	1.00	45.66
	ATOM	1142	O	LEU	144	-12.741	87.571	-36.669	1.00	46.37
	ATOM	1143	OXT	LEU	144	-12.813	89.763	-36.525	1.00	50.66
25	END					-119.496	62.638	-15.481	0.00	0.00

TABLE 6

111									
	ATOM	1	CB	VAL	1	14.586	104.358	-24.172	1.00 69.28
5	ATOM	2	CG1	VAL	1	13.060	104.488	-23.969	1.00 68.26
	ATOM	3	CG2	VAL	1	15.061	105.199	-25.362	1.00 71.75
	ATOM	4	C	VAL	1	15.141	103.681	-21.818	1.00 64.86
	ATOM	5	O	VAL	1	14.357	103.845	-20.872	1.00 64.02
	ATOM	6	N	VAL	1	16.794	105.012	-23.137	1.00 65.10
10	ATOM	7	CA	VAL	1	15.340	104.787	-22.872	1.00 65.98
	ATOM	8	N	THR	2	15.840	102.556	-21.990	1.00 61.42
	ATOM	9	CA	THR	2	15.732	101.442	-21.048	1.00 56.64
	ATOM	10	CB	THR	2	15.231	100.157	-21.731	1.00 56.77
	ATOM	11	OG1	THR	2	16.089	99.849	-22.834	1.00 57.45
15	ATOM	12	CG2	THR	2	13.800	100.323	-22.229	1.00 55.68
	ATOM	13	C	THR	2	17.070	101.114	-20.414	1.00 53.08
	ATOM	14	O	THR	2	18.111	101.603	-20.840	1.00 52.00
	ATOM	15	N	GLN	3	17.033	100.261	-19.398	1.00 50.34
	ATOM	16	CA	GLN	3	18.244	99.856	-18.697	1.00 45.97
20	ATOM	17	CB	GLN	3	18.077	100.074	-17.197	1.00 46.09
	ATOM	18	CG	GLN	3	17.549	101.437	-16.833	1.00 49.60
	ATOM	19	CD	GLN	3	17.478	101.634	-15.340	1.00 50.43
	ATOM	20	OE1	GLN	3	18.505	101.698	-14.673	1.00 50.94
	ATOM	21	NE2	GLN	3	16.262	101.723	-14.805	1.00 49.99
25	ATOM	22	C	GLN	3	18.558	98.395	-18.956	1.00 42.41
	ATOM	23	O	GLN	3	17.860	97.501	-18.483	1.00 37.34
	ATOM	24	N	ASP	4	19.614	98.153	-19.715	1.00 40.32
	ATOM	25	CA	ASP	4	20.011	96.792	-20.027	1.00 39.37
	ATOM	26	CB	ASP	4	21.116	96.795	-21.087	1.00 44.56
30	ATOM	27	CG	ASP	4	20.639	97.328	-22.422	1.00 45.76
	ATOM	28	OD1	ASP	4	19.596	98.021	-22.440	1.00 48.58
	ATOM	29	OD2	ASP	4	21.307	97.062	-23.444	1.00 45.55
	ATOM	30	C	ASP	4	20.519	96.124	-18.768	1.00 36.30
	ATOM	31	O	ASP	4	21.104	96.770	-17.903	1.00 36.29
35	ATOM	32	N	CYS	5	20.280	94.826	-18.663	1.00 35.11
	ATOM	33	CA	CYS	5	20.748	94.062	-17.521	1.00 35.61
	ATOM	34	CB	CYS	5	19.873	94.317	-16.290	1.00 34.96
	ATOM	35	SG	CYS	5	18.119	94.367	-16.599	1.00 35.34
	ATOM	36	C	CYS	5	20.772	92.589	-17.870	1.00 31.80
40	ATOM	37	O	CYS	5	20.014	92.129	-18.715	1.00 31.85
	ATOM	38	N	LEU	6	21.681	91.863	-17.238	1.00 27.72
	ATOM	39	CA	LEU	6	21.818	90.438	-17.460	1.00 30.80
	ATOM	40	CB	LEU	6	22.989	90.160	-18.394	1.00 29.25
	ATOM	41	CG	LEU	6	23.335	88.698	-18.685	1.00 29.47
45	ATOM	42	CD1	LEU	6	24.038	88.598	-20.012	1.00 30.65
	ATOM	43	CD2	LEU	6	24.218	88.159	-17.599	1.00 35.27
	ATOM	44	C	LEU	6	22.054	89.796	-16.107	1.00 31.83
	ATOM	45	O	LEU	6	22.936	90.218	-15.369	1.00 33.45
	ATOM	46	N	GLN	7	21.260	88.785	-15.774	1.00 32.23
50	ATOM	47	CA	GLN	7	21.398	88.120	-14.497	1.00 30.26
	ATOM	48	CB	GLN	7	20.111	88.272	-13.698	1.00 29.56
	ATOM	49	CG	GLN	7	20.215	87.884	-12.239	1.00 25.09
	ATOM	50	CD	GLN	7	18.985	88.301	-11.459	1.00 26.50
	ATOM	51	OE1	GLN	7	17.904	87.735	-11.619	1.00 24.40
55	ATOM	52	NE2	GLN	7	19.140	89.310	-10.621	1.00 27.03
	ATOM	53	C	GLN	7	21.726	86.657	-14.700	1.00 32.11
	ATOM	54	O	GLN	7	21.203	86.011	-15.600	1.00 34.24
	ATOM	55	N	LEU	8	22.615	86.150	-13.859	1.00 33.73
	ATOM	56	CA	LEU	8	23.042	84.761	-13.912	1.00 35.41
60	ATOM	57	CB	LEU	8	24.568	84.687	-14.093	1.00 32.44
	ATOM	58	CG	LEU	8	25.188	84.580	-15.494	1.00 27.69
	ATOM	59	CD1	LEU	8	24.221	85.009	-16.567	1.00 26.43
	ATOM	60	CD2	LEU	8	26.445	85.406	-15.521	1.00 18.64
	ATOM	61	C	LEU	8	22.625	84.037	-12.641	1.00 36.87
65	ATOM	62	O	LEU	8	22.579	84.618	-11.558	1.00 36.05
	ATOM	63	N	ILE	9	22.331	82.754	-12.789	1.00 39.25

	ATOM	64	CA	ILE	9	21.893	81.915	-11.686	1.00	35.57
	ATOM	65	CB	ILE	9	20.413	81.522	-11.920	1.00	34.28
	ATOM	66	CG2	ILE	9	20.179	80.052	-11.679	1.00	38.20
	ATOM	67	CG1	ILE	9	19.522	82.362	-11.035	1.00	35.22
5	ATOM	68	CD1	ILE	9	18.088	81.979	-11.180	1.00	43.63
	ATOM	69	C	ILE	9	22.791	80.677	-11.597	1.00	35.28
	ATOM	70	O	ILE	9	23.277	80.185	-12.611	1.00	33.75
	ATOM	71	N	ALA	10	23.026	80.182	-10.388	1.00	35.78
	ATOM	72	CA	ALA	10	23.861	79.000	-10.231	1.00	35.28
10	ATOM	73	CB	ALA	10	24.074	78.700	-8.765	1.00	28.47
	ATOM	74	C	ALA	10	23.222	77.801	-10.919	1.00	36.49
	ATOM	75	O	ALA	10	22.013	77.601	-10.852	1.00	37.41
	ATOM	76	N	ASP	11	24.045	77.004	-11.585	1.00	39.44
	ATOM	77	CA	ASP	11	23.565	75.816	-12.281	1.00	40.77
15	ATOM	78	CB	ASP	11	24.169	75.753	-13.685	1.00	38.90
	ATOM	79	CG	ASP	11	23.813	74.477	-14.408	1.00	40.34
	ATOM	80	OD1	ASP	11	22.760	73.876	-14.086	1.00	39.50
	ATOM	81	OD2	ASP	11	24.583	74.085	-15.304	1.00	37.38
	ATOM	82	C	ASP	11	23.931	74.564	-11.492	1.00	42.25
20	ATOM	83	O	ASP	11	25.026	74.016	-11.633	1.00	40.54
	ATOM	84	N	SER	12	22.995	74.117	-10.664	1.00	42.92
	ATOM	85	CA	SER	12	23.194	72.953	-9.804	1.00	43.84
	ATOM	86	CB	SER	12	21.993	72.789	-8.880	1.00	42.13
	ATOM	87	OG	SER	12	20.804	72.648	-9.636	1.00	43.31
25	ATOM	88	C	SER	12	23.434	71.642	-10.533	1.00	46.10
	ATOM	89	O	SER	12	23.702	70.618	-9.904	1.00	43.90
	ATOM	90	N	GLU	13	23.345	71.671	-11.858	1.00	46.72
	ATOM	91	CA	GLU	13	23.542	70.463	-12.635	1.00	44.93
	ATOM	92	CB	GLU	13	22.451	70.341	-13.684	1.00	46.71
30	ATOM	93	CG	GLU	13	21.103	70.113	-13.058	1.00	58.25
	ATOM	94	CD	GLU	13	20.145	69.416	-14.003	1.00	66.70
	ATOM	95	OE1	GLU	13	20.490	68.291	-14.460	1.00	74.10
	ATOM	96	OE2	GLU	13	19.055	69.982	-14.288	1.00	71.55
	ATOM	97	C	GLU	13	24.917	70.290	-13.273	1.00	43.24
35	ATOM	98	O	GLU	13	25.101	69.432	-14.139	1.00	44.25
	ATOM	99	N	THR	14	25.878	71.103	-12.862	1.00	37.06
	ATOM	100	CA	THR	14	27.237	70.949	-13.368	1.00	36.77
	ATOM	101	CB	THR	14	27.559	71.909	-14.552	1.00	33.39
	ATOM	102	OG1	THR	14	27.743	73.238	-14.068	1.00	41.69
40	ATOM	103	CG2	THR	14	26.434	71.896	-15.574	1.00	31.66
	ATOM	104	C	THR	14	28.167	71.223	-12.191	1.00	34.96
	ATOM	105	O	THR	14	27.833	71.982	-11.290	1.00	35.04
	ATOM	106	N	PRO	15	29.338	70.587	-12.178	1.00	37.11
	ATOM	107	CD	PRO	15	29.821	69.635	-13.190	1.00	36.50
45	ATOM	108	CA	PRO	15	30.323	70.750	-11.102	1.00	38.81
	ATOM	109	CB	PRO	15	31.410	69.742	-11.473	1.00	39.15
	ATOM	110	CG	PRO	15	30.702	68.754	-12.368	1.00	37.32
	ATOM	111	C	PRO	15	30.885	72.163	-11.021	1.00	37.78
	ATOM	112	O	PRO	15	30.947	72.862	-12.028	1.00	35.91
50	ATOM	113	N	THR	16	31.300	72.586	-9.832	1.00	36.67
	ATOM	114	CA	THR	16	31.873	73.916	-9.702	1.00	38.46
	ATOM	115	CB	THR	16	32.128	74.298	-8.231	1.00	38.61
	ATOM	116	OG1	THR	16	33.101	73.413	-7.666	1.00	42.44
	ATOM	117	CG2	THR	16	30.839	74.208	-7.430	1.00	40.11
55	ATOM	118	C	THR	16	33.202	73.863	-10.424	1.00	35.97
	ATOM	119	O	THR	16	33.926	72.883	-10.301	1.00	39.85
	ATOM	120	N	ILE	17	33.521	74.905	-11.180	1.00	33.66
	ATOM	121	CA	ILE	17	34.772	74.941	-11.924	1.00	34.76
	ATOM	122	CB	ILE	17	34.713	75.999	-13.033	1.00	32.44
60	ATOM	123	CG2	ILE	17	36.060	76.112	-13.719	1.00	28.66
	ATOM	124	CG1	ILE	17	33.607	75.633	-14.029	1.00	33.15
	ATOM	125	CD1	ILE	17	33.396	76.649	-15.134	1.00	31.19
	ATOM	126	C	ILE	17	35.989	75.215	-11.050	1.00	40.64
	ATOM	127	O	ILE	17	36.017	76.178	-10.295	1.00	44.07
65	ATOM	128	N	GLN	18	37.001	74.358	-11.155	1.00	45.91
	ATOM	129	CA	GLN	18	38.229	74.521	-10.381	1.00	50.22

	ATOM	130	CB	GLN	18	38.660	73.194	-9.765	1.00	53.09
	ATOM	131	CG	GLN	18	38.793	73.265	-8.267	1.00	57.60
	ATOM	132	CD	GLN	18	37.447	73.441	-7.612	1.00	61.39
	ATOM	133	OE1	GLN	18	36.616	72.529	-7.623	1.00	68.36
5	ATOM	134	NE2	GLN	18	37.212	74.621	-7.055	1.00	64.93
	ATOM	135	C	GLN	18	39.353	75.033	-11.263	1.00	51.56
	ATOM	136	O	GLN	18	39.584	74.516	-12.352	1.00	56.44
	ATOM	137	N	LYS	19	40.060	76.046	-10.790	1.00	52.60
	ATOM	138	CA	LYS	19	41.157	76.594	-11.571	1.00	56.10
10	ATOM	139	CB	LYS	19	40.625	77.322	-12.806	1.00	55.55
	ATOM	140	CG	LYS	19	41.733	77.998	-13.581	1.00	61.15
	ATOM	141	CD	LYS	19	41.279	78.539	-14.920	1.00	64.42
	ATOM	142	CE	LYS	19	42.460	79.192	-15.654	1.00	64.84
	ATOM	143	NZ	LYS	19	42.086	79.718	-17.006	1.00	69.52
15	ATOM	144	C	LYS	19	42.032	77.542	-10.765	1.00	57.41
	ATOM	145	O	LYS	19	41.525	78.404	-10.045	1.00	61.02
	ATOM	146	N	GLY	20	43.348	77.379	-10.902	1.00	58.84
	ATOM	147	CA	GLY	20	44.292	78.220	-10.192	1.00	56.66
	ATOM	148	C	GLY	20	43.980	78.346	-8.713	1.00	56.87
20	ATOM	149	O	GLY	20	44.151	79.422	-8.138	1.00	58.64
	ATOM	150	N	SER	21	43.527	77.249	-8.102	1.00	54.49
	ATOM	151	CA	SER	21	43.180	77.220	-6.679	1.00	53.68
	ATOM	152	CB	SER	21	44.411	77.541	-5.821	1.00	55.80
	ATOM	153	OG	SER	21	44.598	78.936	-5.670	1.00	65.34
25	ATOM	154	C	SER	21	42.016	78.178	-6.330	1.00	51.41
	ATOM	155	O	SER	21	41.892	78.666	-5.193	1.00	47.74
	ATOM	156	N	TYR	22	41.172	78.438	-7.328	1.00	46.25
	ATOM	157	CA	TYR	22	40.000	79.289	-7.178	1.00	40.57
	ATOM	158	CB	TYR	22	40.092	80.504	-8.105	1.00	43.27
30	ATOM	159	CG	TYR	22	40.821	81.681	-7.511	1.00	49.18
	ATOM	160	CD1	TYR	22	42.143	81.572	-7.097	1.00	52.61
	ATOM	161	CE1	TYR	22	42.819	82.658	-6.534	1.00	51.92
	ATOM	162	CD2	TYR	22	40.185	82.908	-7.350	1.00	50.01
	ATOM	163	CE2	TYR	22	40.851	83.999	-6.792	1.00	50.94
35	ATOM	164	CZ	TYR	22	42.166	83.867	-6.385	1.00	50.78
	ATOM	165	OH	TYR	22	42.822	84.943	-5.830	1.00	53.87
	ATOM	166	C	TYR	22	38.777	78.462	-7.559	1.00	37.85
	ATOM	167	O	TYR	22	38.868	77.566	-8.397	1.00	38.70
	ATOM	168	N	THR	23	37.639	78.740	-6.936	1.00	33.81
40	ATOM	169	CA	THR	23	36.417	78.016	-7.267	1.00	32.40
	ATOM	170	CB	THR	23	35.657	77.518	-6.012	1.00	31.58
	ATOM	171	OG1	THR	23	36.512	76.692	-5.214	1.00	30.76
	ATOM	172	CG2	THR	23	34.441	76.717	-6.424	1.00	24.08
	ATOM	173	C	THR	23	35.504	78.970	-8.017	1.00	33.28
45	ATOM	174	O	THR	23	35.298	80.100	-7.589	1.00	30.44
	ATOM	175	N	PHE	24	34.964	78.513	-9.137	1.00	32.16
	ATOM	176	CA	PHE	24	34.067	79.335	-9.930	1.00	30.68
	ATOM	177	CB	PHE	24	34.638	79.559	-11.329	1.00	28.20
	ATOM	178	CG	PHE	24	35.893	80.372	-11.343	1.00	31.00
50	ATOM	179	CD1	PHE	24	37.124	79.778	-11.123	1.00	31.69
	ATOM	180	CD2	PHE	24	35.842	81.749	-11.545	1.00	31.42
	ATOM	181	CE1	PHE	24	38.285	80.541	-11.103	1.00	32.57
	ATOM	182	CE2	PHE	24	36.995	82.515	-11.527	1.00	25.01
	ATOM	183	CZ	PHE	24	38.218	81.909	-11.305	1.00	30.12
55	ATOM	184	C	PHE	24	32.679	78.715	-10.035	1.00	31.73
	ATOM	185	O	PHE	24	32.531	77.542	-10.363	1.00	33.31
	ATOM	186	N	VAL	25	31.664	79.513	-9.746	1.00	32.92
	ATOM	187	CA	VAL	25	30.296	79.050	-9.820	1.00	30.84
	ATOM	188	CB	VAL	25	29.331	80.088	-9.226	1.00	30.29
60	ATOM	189	CG1	VAL	25	27.907	79.605	-9.351	1.00	28.13
	ATOM	190	CG2	VAL	25	29.677	80.349	-7.786	1.00	29.53
	ATOM	191	C	VAL	25	29.901	78.818	-11.274	1.00	36.08
	ATOM	192	O	VAL	25	30.220	79.621	-12.148	1.00	35.82
	ATOM	193	N	PRO	26	29.223	77.696	-11.556	1.00	38.03
65	ATOM	194	CD	PRO	26	29.063	76.539	-10.662	1.00	38.28
	ATOM	195	CA	PRO	26	28.776	77.361	-12.916	1.00	38.08

	ATOM	196	CB	PRO	26	28.474	75.863	-12.829	1.00	37.62
	ATOM	197	CG	PRO	26	29.221	75.401	-11.619	1.00	36.31
	ATOM	198	C	PRO	26	27.498	78.165	-13.148	1.00	40.85
	ATOM	199	O	PRO	26	26.475	77.897	-12.518	1.00	42.27
5	ATOM	200	N	TRP	27	27.536	79.144	-14.040	1.00	40.86
	ATOM	201	CA	TRP	27	26.348	79.962	-14.253	1.00	37.05
	ATOM	202	CB	TRP	27	26.745	81.398	-14.602	1.00	33.29
	ATOM	203	CG	TRP	27	27.538	82.068	-13.539	1.00	31.06
	ATOM	204	CD2	TRP	27	27.124	82.333	-12.198	1.00	27.79
10	ATOM	205	CE2	TRP	27	28.204	82.948	-11.544	1.00	29.28
	ATOM	206	CE3	TRP	27	25.945	82.110	-11.485	1.00	27.06
	ATOM	207	CD1	TRP	27	28.811	82.522	-13.642	1.00	31.07
	ATOM	208	NE1	TRP	27	29.222	83.052	-12.452	1.00	32.70
	ATOM	209	CZ2	TRP	27	28.145	83.345	-10.205	1.00	25.99
15	ATOM	210	CZ3	TRP	27	25.886	82.504	-10.153	1.00	26.93
	ATOM	211	CH2	TRP	27	26.981	83.114	-9.529	1.00	28.08
	ATOM	212	C	TRP	27	25.379	79.454	-15.296	1.00	38.50
	ATOM	213	O	TRP	27	25.734	78.680	-16.179	1.00	39.07
	ATOM	214	N	LEU	28	24.139	79.913	-15.163	1.00	40.19
20	ATOM	215	CA	LEU	28	23.055	79.580	-16.072	1.00	39.45
	ATOM	216	CB	LEU	28	22.112	78.573	-15.425	1.00	44.18
	ATOM	217	CG	LEU	28	21.163	77.845	-16.375	1.00	46.83
	ATOM	218	CD1	LEU	28	21.985	76.928	-17.286	1.00	45.15
	ATOM	219	CD2	LEU	28	20.140	77.039	-15.571	1.00	49.15
25	ATOM	220	C	LEU	28	22.325	80.901	-16.270	1.00	39.80
	ATOM	221	O	LEU	28	22.145	81.648	-15.318	1.00	42.40
	ATOM	222	N	LEU	29	21.913	81.206	-17.489	1.00	33.58
	ATOM	223	CA	LEU	29	21.222	82.461	-17.714	1.00	29.21
	ATOM	224	CB	LEU	29	20.975	82.688	-19.205	1.00	27.16
30	ATOM	225	CG	LEU	29	20.189	83.964	-19.521	1.00	22.36
	ATOM	226	CD1	LEU	29	21.083	85.177	-19.322	1.00	21.47
	ATOM	227	CD2	LEU	29	19.662	83.919	-20.930	1.00	14.53
	ATOM	228	C	LEU	29	19.887	82.533	-16.982	1.00	29.98
	ATOM	229	O	LEU	29	19.041	81.653	-17.110	1.00	30.36
35	ATOM	230	N	SER	30	19.715	83.595	-16.208	1.00	29.88
	ATOM	231	CA	SER	30	18.476	83.820	-15.487	1.00	28.49
	ATOM	232	CB	SER	30	18.727	84.620	-14.209	1.00	26.13
	ATOM	233	OG	SER	30	17.512	84.917	-13.560	1.00	20.25
	ATOM	234	C	SER	30	17.618	84.621	-16.457	1.00	31.61
40	ATOM	235	O	SER	30	16.485	84.253	-16.752	1.00	35.14
	ATOM	236	N	PHE	31	18.174	85.718	-16.958	1.00	32.12
	ATOM	237	CA	PHE	31	17.479	86.555	-17.921	1.00	30.76
	ATOM	238	CB	PHE	31	16.276	87.251	-17.272	1.00	27.55
	ATOM	239	CG	PHE	31	16.598	88.569	-16.624	1.00	31.23
45	ATOM	240	CD1	PHE	31	16.672	89.732	-17.382	1.00	32.00
	ATOM	241	CD2	PHE	31	16.841	88.647	-15.255	1.00	27.89
	ATOM	242	CE1	PHE	31	16.981	90.946	-16.790	1.00	29.70
	ATOM	243	CE2	PHE	31	17.150	89.858	-14.659	1.00	26.15
	ATOM	244	CZ	PHE	31	17.221	91.009	-15.425	1.00	28.30
50	ATOM	245	C	PHE	31	18.436	87.584	-18.497	1.00	32.06
	ATOM	246	O	PHE	31	19.397	87.998	-17.848	1.00	32.26
	ATOM	247	N	LYS	32	18.174	87.981	-19.730	1.00	34.83
	ATOM	248	CA	LYS	32	18.992	88.973	-20.397	1.00	37.13
	ATOM	249	CB	LYS	32	19.799	88.321	-21.518	1.00	36.55
55	ATOM	250	CG	LYS	32	20.626	89.298	-22.305	1.00	40.62
	ATOM	251	CD	LYS	32	21.184	88.684	-23.564	1.00	44.56
	ATOM	252	CE	LYS	32	21.746	89.791	-24.459	1.00	46.29
	ATOM	253	NZ	LYS	32	22.433	89.269	-25.677	1.00	50.58
	ATOM	254	C	LYS	32	18.039	90.023	-20.963	1.00	36.39
60	ATOM	255	O	LYS	32	17.022	89.684	-21.569	1.00	38.69
	ATOM	256	N	ARG	33	18.357	91.294	-20.747	1.00	37.90
	ATOM	257	CA	ARG	33	17.511	92.374	-21.236	1.00	35.74
	ATOM	258	CB	ARG	33	16.743	92.978	-20.066	1.00	36.79
	ATOM	259	CG	ARG	33	15.874	94.169	-20.402	1.00	31.09
65	ATOM	260	CD	ARG	33	14.968	94.490	-19.220	1.00	31.62
	ATOM	261	NE	ARG	33	13.987	95.533	-19.508	1.00	38.16

	ATOM	262	CZ	ARG	33	14.111	96.800	-19.131	1.00	39.15
	ATOM	263	NH1	ARG	33	15.176	97.187	-18.443	1.00	43.41
	ATOM	264	NH2	ARG	33	13.175	97.681	-19.440	1.00	40.65
	ATOM	265	C	ARG	33	18.362	93.437	-21.922	1.00	37.71
5	ATOM	266	O	ARG	33	19.244	94.037	-21.301	1.00	41.16
	ATOM	267	N	GLY	34	18.111	93.669	-23.205	1.00	36.59
	ATOM	268	CA	GLY	34	18.883	94.666	-23.916	1.00	36.72
	ATOM	269	C	GLY	34	20.092	94.081	-24.615	1.00	40.01
	ATOM	270	O	GLY	34	20.215	92.866	-24.765	1.00	39.85
10	ATOM	271	N	SER	35	21.002	94.956	-25.020	1.00	38.74
	ATOM	272	CA	SER	35	22.196	94.541	-25.741	1.00	40.23
	ATOM	273	CB	SER	35	22.192	95.209	-27.110	1.00	43.02
	ATOM	274	OG	SER	35	22.043	96.621	-26.970	1.00	48.05
	ATOM	275	C	SER	35	23.532	94.841	-25.057	1.00	40.54
15	ATOM	276	O	SER	35	24.551	94.259	-25.425	1.00	43.74
	ATOM	277	N	ALA	36	23.530	95.737	-24.073	1.00	35.56
	ATOM	278	CA	ALA	36	24.753	96.130	-23.382	1.00	29.62
	ATOM	279	CB	ALA	36	24.439	97.232	-22.391	1.00	25.47
	ATOM	280	C	ALA	36	25.530	95.017	-22.685	1.00	31.23
20	ATOM	281	O	ALA	36	26.734	95.136	-22.495	1.00	29.56
	ATOM	282	N	LEU	37	24.855	93.937	-22.305	1.00	34.09
	ATOM	283	CA	LEU	37	25.516	92.841	-21.604	1.00	32.32
	ATOM	284	CB	LEU	37	25.099	92.842	-20.135	1.00	26.47
	ATOM	285	CG	LEU	37	25.435	94.121	-19.372	1.00	25.72
25	ATOM	286	CD1	LEU	37	24.600	94.221	-18.120	1.00	24.61
	ATOM	287	CD2	LEU	37	26.910	94.145	-19.047	1.00	27.53
	ATOM	288	C	LEU	37	25.222	91.481	-22.212	1.00	35.16
	ATOM	289	O	LEU	37	24.112	91.210	-22.655	1.00	36.47
	ATOM	290	N	GLU	38	26.232	90.625	-22.220	1.00	38.30
30	ATOM	291	CA	GLU	38	26.103	89.283	-22.765	1.00	41.17
	ATOM	292	CB	GLU	38	26.748	89.206	-24.143	1.00	46.15
	ATOM	293	CG	GLU	38	25.848	89.553	-25.305	1.00	50.01
	ATOM	294	CD	GLU	38	26.639	89.721	-26.596	1.00	51.18
	ATOM	295	OE1	GLU	38	27.587	88.923	-26.816	1.00	48.40
35	ATOM	296	OE2	GLU	38	26.312	90.648	-27.384	1.00	50.23
	ATOM	297	C	GLU	38	26.790	88.264	-21.876	1.00	42.69
	ATOM	298	O	GLU	38	27.684	88.594	-21.108	1.00	42.55
	ATOM	299	N	GLU	39	26.371	87.014	-21.983	1.00	44.75
	ATOM	300	CA	GLU	39	27.002	85.966	-21.199	1.00	46.50
40	ATOM	301	CB	GLU	39	25.980	84.908	-20.795	1.00	47.52
	ATOM	302	CG	GLU	39	24.973	84.614	-21.886	1.00	58.50
	ATOM	303	CD	GLU	39	24.049	83.466	-21.530	1.00	62.35
	ATOM	304	OE1	GLU	39	23.048	83.254	-22.262	1.00	64.58
	ATOM	305	OE2	GLU	39	24.330	82.772	-20.524	1.00	63.04
45	ATOM	306	C	GLU	39	28.034	85.374	-22.142	1.00	44.56
	ATOM	307	O	GLU	39	27.762	85.185	-23.320	1.00	48.11
	ATOM	308	N	LYS	40	29.226	85.103	-21.635	1.00	40.71
	ATOM	309	CA	LYS	40	30.277	84.549	-22.465	1.00	38.08
	ATOM	310	CB	LYS	40	31.057	85.672	-23.146	1.00	41.28
50	ATOM	311	CG	LYS	40	32.294	85.200	-23.899	1.00	41.52
	ATOM	312	CD	LYS	40	33.192	86.375	-24.262	1.00	42.73
	ATOM	313	CE	LYS	40	34.508	85.924	-24.871	1.00	41.97
	ATOM	314	NZ	LYS	40	35.416	87.090	-25.100	1.00	45.94
	ATOM	315	C	LYS	40	31.230	83.703	-21.648	1.00	39.66
55	ATOM	316	O	LYS	40	31.947	84.209	-20.788	1.00	36.18
	ATOM	317	N	GLU	41	31.228	82.406	-21.926	1.00	40.16
	ATOM	318	CA	GLU	41	32.105	81.466	-21.245	1.00	42.80
	ATOM	319	CB	GLU	41	33.518	81.625	-21.793	1.00	45.31
	ATOM	320	CG	GLU	41	33.558	81.474	-23.309	1.00	56.07
60	ATOM	321	CD	GLU	41	34.829	82.035	-23.941	1.00	59.42
	ATOM	322	OE1	GLU	41	35.150	83.227	-23.709	1.00	63.71
	ATOM	323	OE2	GLU	41	35.504	81.283	-24.678	1.00	63.71
	ATOM	324	C	GLU	41	32.086	81.643	-19.735	1.00	38.93
	ATOM	325	O	GLU	41	33.127	81.811	-19.106	1.00	39.03
65	ATOM	326	N	ASN	42	30.886	81.621	-19.172	1.00	34.69
	ATOM	327	CA	ASN	42	30.689	81.743	-17.738	1.00	32.96

	ATOM	328	CB	ASN	42	31.499	80.679	-17.007	1.00	32.65
	ATOM	329	CG	ASN	42	30.801	80.179	-15.781	1.00	35.50
	ATOM	330	OD1	ASN	42	31.410	80.017	-14.732	1.00	40.26
	ATOM	331	ND2	ASN	42	29.508	79.921	-15.906	1.00	29.28
5	ATOM	332	C	ASN	42	31.021	83.104	-17.146	1.00	31.44
	ATOM	333	O	ASN	42	31.199	83.233	-15.939	1.00	22.78
	ATOM	334	N	LYS	43	31.095	84.116	-17.996	1.00	33.26
	ATOM	335	CA	LYS	43	31.404	85.463	-17.546	1.00	33.68
	ATOM	336	CB	LYS	43	32.837	85.832	-17.926	1.00	33.55
10	ATOM	337	CG	LYS	43	33.887	85.037	-17.191	1.00	40.49
	ATOM	338	CD	LYS	43	35.269	85.383	-17.682	1.00	45.39
	ATOM	339	CE	LYS	43	35.448	84.960	-19.127	1.00	52.57
	ATOM	340	NZ	LYS	43	36.839	85.198	-19.590	1.00	62.06
	ATOM	341	C	LYS	43	30.448	86.443	-18.187	1.00	32.05
15	ATOM	342	O	LYS	43	29.752	86.102	-19.134	1.00	37.32
	ATOM	343	N	ILE	44	30.399	87.659	-17.666	1.00	31.31
	ATOM	344	CA	ILE	44	29.525	88.656	-18.250	1.00	29.52
	ATOM	345	CB	ILE	44	28.806	89.471	-17.173	1.00	26.38
	ATOM	346	CG2	ILE	44	27.941	90.529	-17.825	1.00	26.09
20	ATOM	347	CG1	ILE	44	27.930	88.542	-16.332	1.00	28.61
	ATOM	348	CD1	ILE	44	27.152	89.231	-15.254	1.00	21.51
	ATOM	349	C	ILE	44	30.354	89.573	-19.131	1.00	31.60
	ATOM	350	O	ILE	44	31.317	90.188	-18.677	1.00	34.50
	ATOM	351	N	LEU	45	29.985	89.641	-20.403	1.00	31.44
25	ATOM	352	CA	LEU	45	30.686	90.475	-21.368	1.00	31.40
	ATOM	353	CB	LEU	45	30.811	89.729	-22.694	1.00	33.60
	ATOM	354	CG	LEU	45	31.423	90.515	-23.850	1.00	34.36
	ATOM	355	CD1	LEU	45	32.880	90.817	-23.566	1.00	29.00
	ATOM	356	CD2	LEU	45	31.280	89.705	-25.124	1.00	32.42
30	ATOM	357	C	LEU	45	29.984	91.810	-21.597	1.00	27.80
	ATOM	358	O	LEU	45	28.787	91.860	-21.859	1.00	27.94
	ATOM	359	N	VAL	46	30.747	92.891	-21.495	1.00	28.00
	ATOM	360	CA	VAL	46	30.222	94.233	-21.701	1.00	31.53
	ATOM	361	CB	VAL	46	31.042	95.251	-20.892	1.00	27.89
35	ATOM	362	CG1	VAL	46	30.474	96.639	-21.059	1.00	26.10
	ATOM	363	CG2	VAL	46	31.048	94.853	-19.442	1.00	23.39
	ATOM	364	C	VAL	46	30.293	94.587	-23.195	1.00	33.94
	ATOM	365	O	VAL	46	31.364	94.524	-23.802	1.00	39.68
	ATOM	366	N	LYS	47	29.159	94.962	-23.782	1.00	33.59
40	ATOM	367	CA	LYS	47	29.120	95.310	-25.200	1.00	37.05
	ATOM	368	CB	LYS	47	27.977	94.564	-25.897	1.00	34.49
	ATOM	369	CG	LYS	47	27.942	93.077	-25.591	1.00	37.53
	ATOM	370	CD	LYS	47	29.251	92.394	-25.961	1.00	41.69
	ATOM	371	CE	LYS	47	29.185	91.788	-27.351	1.00	42.95
45	ATOM	372	NZ	LYS	47	28.613	92.738	-28.332	1.00	44.50
	ATOM	373	C	LYS	47	28.984	96.812	-25.456	1.00	37.34
	ATOM	374	O	LYS	47	29.151	97.270	-26.582	1.00	43.04
	ATOM	375	N	GLU	48	28.677	97.566	-24.411	1.00	36.99
	ATOM	376	CA	GLU	48	28.538	99.017	-24.506	1.00	37.31
50	ATOM	377	CB	GLU	48	27.076	99.428	-24.473	1.00	36.53
	ATOM	378	CG	GLU	48	26.273	99.050	-25.682	1.00	46.74
	ATOM	379	CD	GLU	48	24.819	99.461	-25.532	1.00	52.39
	ATOM	380	OE1	GLU	48	24.571	100.580	-25.004	1.00	51.33
	ATOM	381	OE2	GLU	48	23.930	98.672	-25.944	1.00	54.55
55	ATOM	382	C	GLU	48	29.209	99.605	-23.290	1.00	36.94
	ATOM	383	O	GLU	48	28.910	99.194	-22.176	1.00	40.21
	ATOM	384	N	THR	49	30.100	100.571	-23.470	1.00	36.34
	ATOM	385	CA	THR	49	30.760	101.124	-22.298	1.00	35.48
	ATOM	386	CB	THR	49	32.090	101.842	-22.662	1.00	31.16
60	ATOM	387	OG1	THR	49	31.864	103.245	-22.775	1.00	32.48
	ATOM	388	CG2	THR	49	32.638	101.323	-23.969	1.00	30.36
	ATOM	389	C	THR	49	29.823	102.080	-21.555	1.00	34.59
	ATOM	390	O	THR	49	28.945	102.700	-22.150	1.00	35.13
	ATOM	391	N	GLY	50	30.010	102.169	-20.240	1.00	32.49
65	ATOM	392	CA	GLY	50	29.185	103.038	-19.430	1.00	28.17
	ATOM	393	C	GLY	50	29.302	102.682	-17.965	1.00	29.99

	ATOM	394	O	GLY	50	30.245	102.019	-17.564	1.00	29.76
	ATOM	395	N	TYR	51	28.344	103.136	-17.165	1.00	31.68
	ATOM	396	CA	TYR	51	28.328	102.848	-15.735	1.00	32.88
	ATOM	397	CB	TYR	51	27.964	104.103	-14.934	1.00	35.97
5	ATOM	398	CG	TYR	51	29.050	105.138	-14.945	1.00	43.03
	ATOM	399	CD1	TYR	51	29.203	106.009	-16.027	1.00	44.15
	ATOM	400	CE1	TYR	51	30.268	106.910	-16.084	1.00	47.81
	ATOM	401	CD2	TYR	51	29.982	105.194	-13.911	1.00	47.08
	ATOM	402	CE2	TYR	51	31.053	106.086	-13.954	1.00	51.79
10	ATOM	403	CZ	TYR	51	31.195	106.940	-15.046	1.00	51.15
	ATOM	404	OH	TYR	51	32.286	107.787	-15.110	1.00	52.20
	ATOM	405	C	TYR	51	27.341	101.732	-15.419	1.00	31.97
	ATOM	406	O	TYR	51	26.171	101.796	-15.788	1.00	30.01
	ATOM	407	N	PHE	52	27.822	100.702	-14.731	1.00	31.50
15	ATOM	408	CA	PHE	52	26.972	99.575	-14.387	1.00	30.41
	ATOM	409	CB	PHE	52	27.473	98.299	-15.068	1.00	31.30
	ATOM	410	CG	PHE	52	27.518	98.374	-16.562	1.00	29.70
	ATOM	411	CD1	PHE	52	28.494	99.117	-17.204	1.00	26.20
	ATOM	412	CD2	PHE	52	26.587	97.687	-17.327	1.00	28.38
20	ATOM	413	CE1	PHE	52	28.539	99.175	-18.582	1.00	27.17
	ATOM	414	CE2	PHE	52	26.626	97.741	-18.708	1.00	27.76
	ATOM	415	CZ	PHE	52	27.601	98.484	-19.337	1.00	27.27
	ATOM	416	C	PHE	52	26.894	99.305	-12.899	1.00	27.65
	ATOM	417	O	PHE	52	27.849	99.523	-12.164	1.00	24.85
25	ATOM	418	N	PHE	53	25.734	98.831	-12.470	1.00	27.18
	ATOM	419	CA	PHE	53	25.514	98.446	-11.085	1.00	27.05
	ATOM	420	CB	PHE	53	24.079	98.733	-10.661	1.00	28.76
	ATOM	421	CG	PHE	53	23.731	98.187	-9.314	1.00	28.00
	ATOM	422	CD1	PHE	53	24.261	98.746	-8.161	1.00	26.49
30	ATOM	423	CD2	PHE	53	22.884	97.094	-9.195	1.00	28.04
	ATOM	424	CE1	PHE	53	23.950	98.227	-6.910	1.00	26.54
	ATOM	425	CE2	PHE	53	22.571	96.571	-7.943	1.00	25.78
	ATOM	426	CZ	PHE	53	23.106	97.140	-6.803	1.00	24.39
	ATOM	427	C	PHE	53	25.741	96.938	-11.132	1.00	27.56
35	ATOM	428	O	PHE	53	25.057	96.233	-11.869	1.00	25.39
	ATOM	429	N	ILE	54	26.708	96.451	-10.362	1.00	28.87
	ATOM	430	CA	ILE	54	27.042	95.031	-10.354	1.00	28.95
	ATOM	431	CB	ILE	54	28.526	94.843	-10.729	1.00	29.60
	ATOM	432	CG2	ILE	54	28.813	93.374	-10.997	1.00	27.42
40	ATOM	433	CG1	ILE	54	28.846	95.673	-11.975	1.00	26.41
	ATOM	434	CD1	ILE	54	30.308	95.843	-12.253	1.00	28.70
	ATOM	435	C	ILE	54	26.768	94.399	-8.990	1.00	28.57
	ATOM	436	O	ILE	54	27.081	94.975	-7.953	1.00	29.17
	ATOM	437	N	TYR	55	26.188	93.206	-8.986	1.00	26.97
45	ATOM	438	CA	TYR	55	25.864	92.549	-7.725	1.00	24.22
	ATOM	439	CB	TYR	55	24.407	92.835	-7.359	1.00	21.82
	ATOM	440	CG	TYR	55	23.407	92.380	-8.393	1.00	27.68
	ATOM	441	CD1	TYR	55	22.852	91.105	-8.342	1.00	27.01
	ATOM	442	CE1	TYR	55	21.952	90.682	-9.304	1.00	27.44
50	ATOM	443	CD2	TYR	55	23.030	93.218	-9.437	1.00	27.16
	ATOM	444	CE2	TYR	55	22.131	92.800	-10.405	1.00	25.39
	ATOM	445	CZ	TYR	55	21.598	91.534	-10.333	1.00	29.02
	ATOM	446	OH	TYR	55	20.717	91.118	-11.299	1.00	31.82
	ATOM	447	C	TYR	55	26.111	91.056	-7.749	1.00	25.16
55	ATOM	448	O	TYR	55	26.245	90.467	-8.798	1.00	28.47
	ATOM	449	N	GLY	56	26.182	90.446	-6.579	1.00	27.44
	ATOM	450	CA	GLY	56	26.417	89.019	-6.517	1.00	22.07
	ATOM	451	C	GLY	56	26.207	88.464	-5.125	1.00	25.83
	ATOM	452	O	GLY	56	26.471	89.141	-4.136	1.00	26.00
60	ATOM	453	N	GLN	57	25.709	87.237	-5.053	1.00	22.56
	ATOM	454	CA	GLN	57	25.482	86.565	-3.784	1.00	20.91
	ATOM	455	CB	GLN	57	24.029	86.695	-3.325	1.00	17.87
	ATOM	456	CG	GLN	57	23.737	85.887	-2.066	1.00	16.86
	ATOM	457	CD	GLN	57	22.328	86.063	-1.540	1.00	22.06
65	ATOM	458	OE1	GLN	57	21.941	87.151	-1.127	1.00	27.26
	ATOM	459	NE2	GLN	57	21.553	84.985	-1.548	1.00	20.20

	ATOM	460	C	GLN	57	25.822	85.097	-3.936	1.00	24.23
	ATOM	461	O	GLN	57	25.594	84.510	-4.992	1.00	25.16
	ATOM	462	N	VAL	58	26.370	84.518	-2.871	1.00	24.62
	ATOM	463	CA	VAL	58	26.753	83.112	-2.838	1.00	26.68
5	ATOM	464	CB	VAL	58	28.278	82.937	-3.040	1.00	25.55
	ATOM	465	CG1	VAL	58	28.675	81.502	-2.794	1.00	15.69
	ATOM	466	CG2	VAL	58	28.678	83.352	-4.440	1.00	25.77
	ATOM	467	C	VAL	58	26.404	82.572	-1.467	1.00	31.06
	ATOM	468	O	VAL	58	26.587	83.270	-0.479	1.00	27.74
10	ATOM	469	N	LEU	59	25.897	81.340	-1.403	1.00	33.82
	ATOM	470	CA	LEU	59	25.570	80.705	-0.122	1.00	29.85
	ATOM	471	CB	LEU	59	24.278	79.898	-0.212	1.00	29.62
	ATOM	472	CG	LEU	59	23.517	79.637	1.095	1.00	28.11
	ATOM	473	CD1	LEU	59	22.488	78.549	0.862	1.00	26.67
15	ATOM	474	CD2	LEU	59	24.445	79.230	2.201	1.00	22.45
	ATOM	475	C	LEU	59	26.714	79.760	0.236	1.00	30.98
	ATOM	476	O	LEU	59	26.952	78.771	-0.454	1.00	32.52
	ATOM	477	N	TYR	60	27.424	80.071	1.315	1.00	33.01
	ATOM	478	CA	TYR	60	28.540	79.241	1.747	1.00	34.51
20	ATOM	479	CB	TYR	60	29.640	80.112	2.345	1.00	40.10
	ATOM	480	CG	TYR	60	30.205	81.082	1.354	1.00	45.37
	ATOM	481	CD1	TYR	60	29.894	82.441	1.419	1.00	47.13
	ATOM	482	CE1	TYR	60	30.350	83.322	0.449	1.00	50.28
	ATOM	483	CD2	TYR	60	30.990	80.633	0.301	1.00	44.35
25	ATOM	484	CE2	TYR	60	31.451	81.497	-0.672	1.00	48.81
	ATOM	485	CZ	TYR	60	31.125	82.836	-0.597	1.00	51.91
	ATOM	486	OH	TYR	60	31.553	83.678	-1.594	1.00	53.77
	ATOM	487	C	TYR	60	28.140	78.171	2.748	1.00	35.49
	ATOM	488	O	TYR	60	27.595	78.462	3.808	1.00	36.13
30	ATOM	489	N	THR	61	28.416	76.924	2.395	1.00	36.19
	ATOM	490	CA	THR	61	28.111	75.797	3.254	1.00	35.36
	ATOM	491	CB	THR	61	27.125	74.846	2.575	1.00	34.22
	ATOM	492	OG1	THR	61	27.623	74.486	1.284	1.00	34.22
	ATOM	493	CG2	THR	61	25.766	75.506	2.428	1.00	25.45
35	ATOM	494	C	THR	61	29.427	75.081	3.522	1.00	37.82
	ATOM	495	O	THR	61	29.480	73.864	3.673	1.00	43.70
	ATOM	496	N	ASP	62	30.491	75.871	3.571	1.00	37.64
	ATOM	497	CA	ASP	62	31.844	75.398	3.804	1.00	38.08
	ATOM	498	CB	ASP	62	32.747	76.026	2.747	1.00	38.42
40	ATOM	499	CG	ASP	62	34.144	75.474	2.764	1.00	41.25
	ATOM	500	OD1	ASP	62	34.712	75.286	1.662	1.00	38.28
	ATOM	501	OD2	ASP	62	34.668	75.245	3.873	1.00	41.37
	ATOM	502	C	ASP	62	32.246	75.851	5.204	1.00	40.90
	ATOM	503	O	ASP	62	31.901	76.954	5.607	1.00	46.73
45	ATOM	504	N	LYS	63	32.967	75.023	5.956	1.00	42.62
	ATOM	505	CA	LYS	63	33.346	75.424	7.311	1.00	40.92
	ATOM	506	CB	LYS	63	33.159	74.258	8.283	1.00	42.79
	ATOM	507	CG	LYS	63	33.956	73.016	7.950	1.00	45.91
	ATOM	508	CD	LYS	63	33.531	71.854	8.849	1.00	46.22
50	ATOM	509	CE	LYS	63	32.047	71.501	8.632	1.00	49.97
	ATOM	510	NZ	LYS	63	31.498	70.523	9.634	1.00	49.98
	ATOM	511	C	LYS	63	34.754	75.991	7.459	1.00	42.59
	ATOM	512	O	LYS	63	35.291	76.039	8.558	1.00	42.46
	ATOM	513	N	THR	64	35.325	76.457	6.356	1.00	44.06
55	ATOM	514	CA	THR	64	36.669	77.016	6.348	1.00	44.94
	ATOM	515	CB	THR	64	37.168	77.191	4.892	1.00	43.79
	ATOM	516	OG1	THR	64	37.073	75.938	4.207	1.00	46.20
	ATOM	517	CG2	THR	64	38.621	77.649	4.858	1.00	42.80
	ATOM	518	C	THR	64	36.862	78.342	7.103	1.00	45.82
60	ATOM	519	O	THR	64	37.518	79.241	6.607	1.00	53.10
	ATOM	520	N	TYR	65	36.306	78.463	8.298	1.00	45.11
	ATOM	521	CA	TYR	65	36.451	79.664	9.133	1.00	45.72
	ATOM	522	CB	TYR	65	37.853	79.696	9.771	1.00	40.46
	ATOM	523	CG	TYR	65	38.901	80.490	9.024	1.00	41.90
65	ATOM	524	CD1	TYR	65	39.061	81.859	9.243	1.00	42.47
	ATOM	525	CE1	TYR	65	40.028	82.603	8.543	1.00	41.83

	ATOM	526	CD2	TYR	65	39.735	79.874	8.085	1.00	41.65
	ATOM	527	CE2	TYR	65	40.704	80.605	7.376	1.00	40.28
	ATOM	528	CZ	TYR	65	40.841	81.966	7.613	1.00	41.90
	ATOM	529	OH	TYR	65	41.786	82.686	6.924	1.00	43.84
5	ATOM	530	C	TYR	65	36.146	81.028	8.493	1.00	45.31
	ATOM	531	O	TYR	65	35.702	81.958	9.179	1.00	49.74
	ATOM	532	N	ALA	66	36.387	81.158	7.196	1.00	42.32
	ATOM	533	CA	ALA	66	36.134	82.408	6.492	1.00	39.57
	ATOM	534	CB	ALA	66	37.274	83.376	6.717	1.00	35.36
10	ATOM	535	C	ALA	66	35.964	82.137	5.008	1.00	39.85
	ATOM	536	O	ALA	66	36.855	81.585	4.359	1.00	37.60
	ATOM	537	N	MET	67	34.807	82.510	4.478	1.00	41.88
	ATOM	538	CA	MET	67	34.519	82.313	3.065	1.00	41.48
	ATOM	539	CB	MET	67	33.374	81.311	2.873	1.00	39.98
15	ATOM	540	CG	MET	67	33.709	79.875	3.254	1.00	37.31
	ATOM	541	SD	MET	67	35.089	79.201	2.319	1.00	35.46
	ATOM	542	CE	MET	67	34.292	78.730	0.781	1.00	34.06
	ATOM	543	C	MET	67	34.147	83.647	2.440	1.00	39.50
	ATOM	544	O	MET	67	33.879	84.622	3.145	1.00	38.44
20	ATOM	545	N	GLY	68	34.135	83.683	1.114	1.00	39.84
	ATOM	546	CA	GLY	68	33.791	84.906	0.417	1.00	37.61
	ATOM	547	C	GLY	68	34.064	84.811	-1.065	1.00	33.76
	ATOM	548	O	GLY	68	34.643	83.835	-1.530	1.00	35.58
	ATOM	549	N	HIS	69	33.634	85.818	-1.813	1.00	35.07
25	ATOM	550	CA	HIS	69	33.853	85.834	-3.249	1.00	33.70
	ATOM	551	CB	HIS	69	32.578	85.414	-4.006	1.00	37.02
	ATOM	552	CG	HIS	69	31.354	86.196	-3.640	1.00	34.65
	ATOM	553	CD2	HIS	69	30.684	87.162	-4.309	1.00	35.63
	ATOM	554	ND1	HIS	69	30.668	86.003	-2.461	1.00	37.68
30	ATOM	555	CE1	HIS	69	29.628	86.815	-2.418	1.00	32.64
	ATOM	556	NE2	HIS	69	29.614	87.528	-3.529	1.00	35.90
	ATOM	557	C	HIS	69	34.319	87.201	-3.729	1.00	33.98
	ATOM	558	O	HIS	69	34.290	88.169	-2.986	1.00	30.94
	ATOM	559	N	LEU	70	34.755	87.268	-4.979	1.00	35.35
35	ATOM	560	CA	LEU	70	35.229	88.514	-5.563	1.00	30.76
	ATOM	561	CB	LEU	70	36.717	88.419	-5.879	1.00	29.59
	ATOM	562	CG	LEU	70	37.642	87.712	-4.893	1.00	30.71
	ATOM	563	CD1	LEU	70	38.994	87.518	-5.534	1.00	25.60
	ATOM	564	CD2	LEU	70	37.751	88.517	-3.619	1.00	27.39
40	ATOM	565	C	LEU	70	34.503	88.789	-6.876	1.00	33.19
	ATOM	566	O	LEU	70	34.282	87.878	-7.673	1.00	37.07
	ATOM	567	N	ILE	71	34.106	90.037	-7.096	1.00	30.93
	ATOM	568	CA	ILE	71	33.485	90.392	-8.362	1.00	27.94
	ATOM	569	CB	ILE	71	32.324	91.375	-8.188	1.00	28.22
45	ATOM	570	CG2	ILE	71	31.942	91.974	-9.526	1.00	24.54
	ATOM	571	CG1	ILE	71	31.125	90.645	-7.596	1.00	26.34
	ATOM	572	CD1	ILE	71	29.997	91.558	-7.202	1.00	37.50
	ATOM	573	C	ILE	71	34.651	91.072	-9.062	1.00	30.32
	ATOM	574	O	ILE	71	35.125	92.106	-8.605	1.00	23.07
50	ATOM	575	N	GLN	72	35.134	90.477	-10.149	1.00	29.74
	ATOM	576	CA	GLN	72	36.283	91.042	-10.841	1.00	27.85
	ATOM	577	CB	GLN	72	37.395	90.005	-10.891	1.00	26.70
	ATOM	578	CG	GLN	72	37.676	89.399	-9.546	1.00	28.28
	ATOM	579	CD	GLN	72	38.899	88.527	-9.556	1.00	33.11
55	ATOM	580	OE1	GLN	72	39.010	87.614	-10.369	1.00	41.22
	ATOM	581	NE2	GLN	72	39.828	88.796	-8.649	1.00	30.18
	ATOM	582	C	GLN	72	36.020	91.565	-12.237	1.00	29.85
	ATOM	583	O	GLN	72	35.060	91.168	-12.899	1.00	28.20
	ATOM	584	N	ARG	73	36.904	92.454	-12.677	1.00	30.90
60	ATOM	585	CA	ARG	73	36.810	93.056	-13.997	1.00	30.79
	ATOM	586	CB	ARG	73	36.573	94.553	-13.853	1.00	29.18
	ATOM	587	CG	ARG	73	36.548	95.293	-15.156	1.00	28.66
	ATOM	588	CD	ARG	73	36.874	96.740	-14.934	1.00	29.25
	ATOM	589	NE	ARG	73	36.809	97.501	-16.172	1.00	35.78
65	ATOM	590	CZ	ARG	73	37.371	98.688	-16.346	1.00	35.27
	ATOM	591	NH1	ARG	73	38.051	99.252	-15.358	1.00	34.95

	ATOM	592	NH2	ARG	73	37.247	99.308	-17.505	1.00	33.33
	ATOM	593	C	ARG	73	38.077	92.829	-14.818	1.00	32.42
	ATOM	594	O	ARG	73	39.177	93.103	-14.350	1.00	34.19
	ATOM	595	N	LYS	74	37.917	92.310	-16.031	1.00	35.90
5	ATOM	596	CA	LYS	74	39.048	92.095	-16.933	1.00	38.57
	ATOM	597	CB	LYS	74	38.922	90.764	-17.680	1.00	43.52
	ATOM	598	CG	LYS	74	39.181	89.530	-16.833	1.00	52.89
	ATOM	599	CD	LYS	74	38.993	88.220	-17.634	1.00	58.14
	ATOM	600	CE	LYS	74	39.245	86.998	-16.750	1.00	56.75
10	ATOM	601	NZ	LYS	74	39.028	85.715	-17.484	1.00	63.98
	ATOM	602	C	LYS	74	39.026	93.226	-17.954	1.00	40.11
	ATOM	603	O	LYS	74	38.177	93.239	-18.854	1.00	37.55
	ATOM	604	N	LYS	75	39.953	94.168	-17.816	1.00	34.48
	ATOM	605	CA	LYS	75	40.029	95.307	-18.723	1.00	34.24
15	ATOM	606	CB	LYS	75	41.048	96.322	-18.201	1.00	36.16
	ATOM	607	CG	LYS	75	40.814	96.864	-16.792	1.00	35.99
	ATOM	608	CD	LYS	75	41.933	97.849	-16.430	1.00	40.91
	ATOM	609	CE	LYS	75	41.744	98.448	-15.043	1.00	46.16
	ATOM	610	NZ	LYS	75	42.842	99.380	-14.608	1.00	46.03
20	ATOM	611	C	LYS	75	40.442	94.877	-20.134	1.00	30.92
	ATOM	612	O	LYS	75	41.301	94.030	-20.289	1.00	27.47
	ATOM	613	N	VAL	76	39.829	95.472	-21.155	1.00	32.82
	ATOM	614	CA	VAL	76	40.168	95.168	-22.548	1.00	31.51
	ATOM	615	CB	VAL	76	39.109	95.665	-23.549	1.00	29.19
25	ATOM	616	CG1	VAL	76	39.011	94.707	-24.701	1.00	27.09
	ATOM	617	CG2	VAL	76	37.794	95.848	-22.879	1.00	34.94
	ATOM	618	C	VAL	76	41.428	95.937	-22.888	1.00	33.40
	ATOM	619	O	VAL	76	42.312	95.436	-23.574	1.00	32.92
	ATOM	620	N	HIS	77	41.480	97.177	-22.410	1.00	34.88
30	ATOM	621	CA	HIS	77	42.608	98.061	-22.642	1.00	34.59
	ATOM	622	CB	HIS	77	42.112	99.436	-23.099	1.00	35.92
	ATOM	623	CG	HIS	77	41.274	99.397	-24.340	1.00	34.17
	ATOM	624	CD2	HIS	77	40.321	100.241	-24.798	1.00	31.44
	ATOM	625	ND1	HIS	77	41.396	98.406	-25.293	1.00	35.05
35	ATOM	626	CE1	HIS	77	40.556	98.638	-26.283	1.00	29.84
	ATOM	627	NE2	HIS	77	39.894	99.746	-26.008	1.00	38.35
	ATOM	628	C	HIS	77	43.417	98.183	-21.362	1.00	35.63
	ATOM	629	O	HIS	77	42.849	98.289	-20.275	1.00	40.24
	ATOM	630	N	VAL	78	44.742	98.195	-21.491	1.00	37.37
40	ATOM	631	CA	VAL	78	45.590	98.252	-20.314	1.00	41.32
	ATOM	632	CB	VAL	78	46.429	96.962	-20.214	1.00	43.10
	ATOM	633	CG1	VAL	78	47.206	96.948	-18.921	1.00	45.01
	ATOM	634	CG2	VAL	78	45.514	95.751	-20.251	1.00	44.98
	ATOM	635	C	VAL	78	46.489	99.470	-20.064	1.00	43.02
45	ATOM	636	O	VAL	78	46.256	100.199	-19.101	1.00	49.79
	ATOM	637	N	PHE	79	47.510	99.710	-20.876	1.00	41.33
	ATOM	638	CA	PHE	79	48.396	100.874	-20.628	1.00	45.00
	ATOM	639	CB	PHE	79	47.621	102.143	-20.220	1.00	40.51
	ATOM	640	CG	PHE	79	46.555	102.553	-21.180	1.00	39.52
50	ATOM	641	CD1	PHE	79	45.216	102.338	-20.875	1.00	36.80
	ATOM	642	CD2	PHE	79	46.886	103.159	-22.384	1.00	37.89
	ATOM	643	CE1	PHE	79	44.219	102.722	-21.756	1.00	37.52
	ATOM	644	CE2	PHE	79	45.894	103.546	-23.271	1.00	35.59
	ATOM	645	CZ	PHE	79	44.557	103.328	-22.958	1.00	37.58
55	ATOM	646	C	PHE	79	49.430	100.662	-19.523	1.00	43.50
	ATOM	647	O	PHE	79	49.087	100.349	-18.381	1.00	37.85
	ATOM	648	N	GLY	80	50.694	100.877	-19.869	1.00	46.65
	ATOM	649	CA	GLY	80	51.779	100.753	-18.915	1.00	48.38
	ATOM	650	C	GLY	80	51.790	99.532	-18.022	1.00	47.89
60	ATOM	651	O	GLY	80	51.636	98.401	-18.491	1.00	51.15
	ATOM	652	N	ASP	81	51.973	99.766	-16.725	1.00	45.07
	ATOM	653	CA	ASP	81	52.040	98.685	-15.758	1.00	43.99
	ATOM	654	CB	ASP	81	53.136	98.973	-14.724	1.00	45.66
	ATOM	655	CG	ASP	81	52.792	100.132	-13.801	1.00	47.49
65	ATOM	656	OD1	ASP	81	51.845	100.894	-14.099	1.00	51.90
	ATOM	657	OD2	ASP	81	53.479	100.289	-12.773	1.00	48.08

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	ATOM	658	C	ASP	81	50.726	98.395	-15.055	1.00	42.29
	ATOM	659	O	ASP	81	50.719	97.840	-13.957	1.00	39.94
	ATOM	660	N	GLU	82	49.614	98.770	-15.681	1.00	39.91
	ATOM	661	CA	GLU	82	48.299	98.502	-15.103	1.00	36.58
5	ATOM	662	CB	GLU	82	47.173	99.072	-15.966	1.00	34.91
	ATOM	663	CG	GLU	82	46.896	100.537	-15.859	1.00	37.82
	ATOM	664	CD	GLU	82	45.463	100.858	-16.253	1.00	40.75
	ATOM	665	OE1	GLU	82	44.979	100.307	-17.262	1.00	38.15
	ATOM	666	OE2	GLU	82	44.810	101.660	-15.552	1.00	47.99
10	ATOM	667	C	GLU	82	48.073	96.998	-15.048	1.00	36.58
	ATOM	668	O	GLU	82	48.570	96.258	-15.892	1.00	38.40
	ATOM	669	N	LEU	83	47.322	96.551	-14.052	1.00	33.83
	ATOM	670	CA	LEU	83	46.967	95.145	-13.941	1.00	32.70
	ATOM	671	CB	LEU	83	46.698	94.765	-12.485	1.00	30.78
15	ATOM	672	CG	LEU	83	47.856	94.202	-11.665	1.00	29.26
	ATOM	673	CD1	LEU	83	49.123	94.981	-11.917	1.00	29.60
	ATOM	674	CD2	LEU	83	47.486	94.242	-10.205	1.00	28.65
	ATOM	675	C	LEU	83	45.679	95.094	-14.733	1.00	33.88
	ATOM	676	O	LEU	83	44.814	95.943	-14.546	1.00	38.46
20	ATOM	677	N	SER	84	45.543	94.123	-15.625	1.00	34.94
	ATOM	678	CA	SER	84	44.335	94.041	-16.430	1.00	39.77
	ATOM	679	CB	SER	84	44.610	93.282	-17.727	1.00	38.92
	ATOM	680	OG	SER	84	45.079	91.985	-17.450	1.00	43.11
	ATOM	681	C	SER	84	43.159	93.405	-15.694	1.00	39.89
25	ATOM	682	O	SER	84	42.021	93.449	-16.159	1.00	42.52
	ATOM	683	N	LEU	85	43.439	92.818	-14.540	1.00	40.92
	ATOM	684	CA	LEU	85	42.405	92.191	-13.740	1.00	34.66
	ATOM	685	CB	LEU	85	42.812	90.760	-13.389	1.00	35.12
	ATOM	686	CG	LEU	85	41.811	89.764	-12.786	1.00	36.92
30	ATOM	687	CD1	LEU	85	41.432	90.184	-11.387	1.00	38.24
	ATOM	688	CD2	LEU	85	40.582	89.667	-13.677	1.00	36.08
	ATOM	689	C	LEU	85	42.246	93.024	-12.482	1.00	35.30
	ATOM	690	O	LEU	85	43.163	93.137	-11.673	1.00	32.85
	ATOM	691	N	VAL	86	41.080	93.634	-12.334	1.00	33.26
35	ATOM	692	CA	VAL	86	40.795	94.460	-11.172	1.00	34.57
	ATOM	693	CB	VAL	86	40.334	95.890	-11.576	1.00	35.39
	ATOM	694	CG1	VAL	86	39.883	96.663	-10.352	1.00	31.44
	ATOM	695	CG2	VAL	86	41.466	96.627	-12.264	1.00	38.08
	ATOM	696	C	VAL	86	39.679	93.824	-10.377	1.00	32.85
40	ATOM	697	O	VAL	86	38.670	93.409	-10.941	1.00	35.10
	ATOM	698	N	THR	87	39.851	93.730	-9.071	1.00	33.41
	ATOM	699	CA	THR	87	38.785	93.177	-8.271	1.00	33.56
	ATOM	700	CB	THR	87	39.314	92.113	-7.260	1.00	32.70
	ATOM	701	OG1	THR	87	38.650	92.257	-5.998	1.00	34.29
45	ATOM	702	CG2	THR	87	40.807	92.222	-7.100	1.00	34.95
	ATOM	703	C	THR	87	38.011	94.303	-7.574	1.00	30.57
	ATOM	704	O	THR	87	38.555	95.048	-6.762	1.00	28.08
	ATOM	705	N	LEU	88	36.750	94.448	-7.970	1.00	28.42
	ATOM	706	CA	LEU	88	35.848	95.429	-7.397	1.00	31.31
50	ATOM	707	CB	LEU	88	34.803	95.886	-8.424	1.00	28.21
	ATOM	708	CG	LEU	88	35.001	96.435	-9.841	1.00	29.12
	ATOM	709	CD1	LEU	88	36.438	96.694	-10.131	1.00	31.12
	ATOM	710	CD2	LEU	88	34.420	95.456	-10.833	1.00	28.72
	ATOM	711	C	LEU	88	35.089	94.693	-6.280	1.00	41.15
55	ATOM	712	O	LEU	88	34.845	93.486	-6.365	1.00	52.30
	ATOM	713	N	PHE	89	34.727	95.386	-5.217	1.00	39.23
	ATOM	714	CA	PHE	89	33.925	94.733	-4.179	1.00	42.27
	ATOM	715	CB	PHE	89	32.582	94.356	-4.810	1.00	35.63
	ATOM	716	CG	PHE	89	32.049	95.434	-5.717	1.00	35.06
60	ATOM	717	CD1	PHE	89	31.501	95.116	-6.955	1.00	31.56
	ATOM	718	CD2	PHE	89	32.215	96.788	-5.377	1.00	30.96
	ATOM	719	CE1	PHE	89	31.141	96.129	-7.851	1.00	33.76
	ATOM	720	CE2	PHE	89	31.862	97.802	-6.260	1.00	27.38
	ATOM	721	CZ	PHE	89	31.328	97.474	-7.501	1.00	30.29
65	ATOM	722	C	PHE	89	34.535	93.560	-3.379	1.00	39.99
	ATOM	723	O	PHE	89	35.520	93.755	-2.660	1.00	45.34

	ATOM	724	N	ARG	90	33.973	92.363	-3.456	1.00	38.92
	ATOM	725	CA	ARG	90	34.515	91.260	-2.642	1.00	41.66
	ATOM	726	CB	ARG	90	36.058	91.271	-2.652	1.00	39.24
	ATOM	727	CG	ARG	90	36.721	90.873	-1.319	1.00	29.41
5	ATOM	728	CD	ARG	90	38.259	90.876	-1.403	1.00	31.13
	ATOM	729	NE	ARG	90	38.889	90.771	-0.085	1.00	31.75
	ATOM	730	CZ	ARG	90	39.614	91.730	0.496	1.00	28.86
	ATOM	731	NH1	ARG	90	39.830	92.885	-0.111	1.00	25.48
	ATOM	732	NH2	ARG	90	40.104	91.544	1.711	1.00	29.84
10	ATOM	733	C	ARG	90	34.057	91.270	-1.170	1.00	39.23
	ATOM	734	O	ARG	90	34.158	92.287	-0.487	1.00	31.33
	ATOM	735	N	CYS	91	33.554	90.130	-0.690	1.00	40.12
	ATOM	736	CA	CYS	91	33.141	90.025	0.700	1.00	37.09
	ATOM	737	C	CYS	91	33.553	88.778	1.430	1.00	37.21
15	ATOM	738	O	CYS	91	33.989	87.806	0.827	1.00	37.46
	ATOM	739	CB	CYS	91	31.651	90.203	0.872	1.00	40.13
	ATOM	740	SG	CYS	91	30.427	89.079	0.109	1.00	41.64
	ATOM	741	N	ILE	92	33.400	88.826	2.749	1.00	34.02
	ATOM	742	CA	ILE	92	33.788	87.722	3.615	1.00	33.22
20	ATOM	743	CB	ILE	92	35.114	88.053	4.331	1.00	32.88
	ATOM	744	CG2	ILE	92	35.621	86.839	5.080	1.00	32.61
	ATOM	745	CG1	ILE	92	36.156	88.501	3.304	1.00	34.64
	ATOM	746	CD1	ILE	92	37.443	89.040	3.909	1.00	32.30
	ATOM	747	C	ILE	92	32.742	87.402	4.675	1.00	30.78
25	ATOM	748	O	ILE	92	31.975	88.256	5.069	1.00	33.43
	ATOM	749	N	GLN	93	32.704	86.149	5.113	1.00	29.35
	ATOM	750	CA	GLN	93	31.786	85.712	6.152	1.00	28.69
	ATOM	751	CB	GLN	93	30.539	85.045	5.570	1.00	28.17
	ATOM	752	CG	GLN	93	29.393	85.962	5.135	1.00	25.87
30	ATOM	753	CD	GLN	93	28.817	86.798	6.257	1.00	24.22
	ATOM	754	OE1	GLN	93	29.285	87.894	6.521	1.00	28.90
	ATOM	755	NE2	GLN	93	27.796	86.278	6.925	1.00	25.29
	ATOM	756	C	GLN	93	32.515	84.683	6.987	1.00	31.44
	ATOM	757	O	GLN	93	33.048	83.724	6.449	1.00	32.07
35	ATOM	758	N	ASN	94	32.573	84.888	8.295	1.00	35.77
	ATOM	759	CA	ASN	94	33.208	83.906	9.153	1.00	33.56
	ATOM	760	CB	ASN	94	33.384	84.448	10.573	1.00	34.70
	ATOM	761	CG	ASN	94	34.602	85.326	10.715	1.00	33.01
	ATOM	762	OD1	ASN	94	35.696	84.939	10.342	1.00	30.56
40	ATOM	763	ND2	ASN	94	34.419	86.506	11.272	1.00	36.61
	ATOM	764	C	ASN	94	32.255	82.710	9.159	1.00	35.80
	ATOM	765	O	ASN	94	31.034	82.878	9.135	1.00	36.63
	ATOM	766	N	MET	95	32.806	81.504	9.178	1.00	35.73
	ATOM	767	CA	MET	95	31.979	80.303	9.170	1.00	34.73
45	ATOM	768	CB	MET	95	32.425	79.379	8.033	1.00	32.97
	ATOM	769	CG	MET	95	32.460	80.025	6.655	1.00	27.35
	ATOM	770	SD	MET	95	30.902	80.791	6.164	1.00	25.14
	ATOM	771	CE	MET	95	29.872	79.403	5.833	1.00	15.55
	ATOM	772	C	MET	95	32.033	79.545	10.498	1.00	36.36
50	ATOM	773	O	MET	95	33.027	79.623	11.226	1.00	40.70
	ATOM	774	N	PRO	96	30.953	78.825	10.842	1.00	36.06
	ATOM	775	CD	PRO	96	29.618	78.969	10.261	1.00	35.32
	ATOM	776	CA	PRO	96	30.874	78.040	12.074	1.00	40.00
	ATOM	777	CB	PRO	96	29.376	77.945	12.345	1.00	33.97
55	ATOM	778	CG	PRO	96	28.779	78.993	11.493	1.00	37.05
	ATOM	779	C	PRO	96	31.457	76.663	11.766	1.00	46.34
	ATOM	780	O	PRO	96	31.665	76.325	10.601	1.00	47.59
	ATOM	781	N	GLU	97	31.706	75.860	12.795	1.00	52.14
	ATOM	782	CA	GLU	97	32.263	74.529	12.587	1.00	57.27
60	ATOM	783	CB	GLU	97	33.035	74.074	13.825	1.00	62.29
	ATOM	784	CG	GLU	97	34.093	73.007	13.555	1.00	72.31
	ATOM	785	CD	GLU	97	35.505	73.601	13.457	1.00	79.34
	ATOM	786	OE1	GLU	97	35.988	74.188	14.475	1.00	84.94
	ATOM	787	OE2	GLU	97	36.129	73.485	12.365	1.00	81.39
65	ATOM	788	C	GLU	97	31.126	73.549	12.320	1.00	56.67
	ATOM	789	O	GLU	97	31.260	72.623	11.516	1.00	59.18

	ATOM	790	N	THR	98	29.999	73.772	12.986	1.00	57.55
	ATOM	791	CA	THR	98	28.840	72.895	12.869	1.00	59.71
	ATOM	792	CB	THR	98	27.807	73.232	13.936	1.00	59.24
	ATOM	793	OG1	THR	98	27.417	74.610	13.776	1.00	68.81
5	ATOM	794	CG2	THR	98	28.397	72.991	15.341	1.00	56.15
	ATOM	795	C	THR	98	28.120	72.866	11.522	1.00	60.32
	ATOM	796	O	THR	98	28.525	72.124	10.619	1.00	65.03
	ATOM	797	N	LEU	99	27.047	73.648	11.391	1.00	54.94
	ATOM	798	CA	LEU	99	26.263	73.672	10.153	1.00	51.13
10	ATOM	799	CB	LEU	99	24.782	73.497	10.482	1.00	50.56
	ATOM	800	CG	LEU	99	24.338	72.102	10.913	1.00	52.80
	ATOM	801	CD1	LEU	99	22.981	72.169	11.611	1.00	48.38
	ATOM	802	CD2	LEU	99	24.280	71.200	9.689	1.00	50.80
	ATOM	803	C	LEU	99	26.454	74.941	9.317	1.00	50.20
15	ATOM	804	O	LEU	99	25.592	75.832	9.310	1.00	50.50
	ATOM	805	N	PRO	100	27.570	75.024	8.569	1.00	47.97
	ATOM	806	CD	PRO	100	28.562	73.964	8.335	1.00	45.68
	ATOM	807	CA	PRO	100	27.864	76.193	7.737	1.00	46.54
	ATOM	808	CB	PRO	100	29.120	75.771	6.972	1.00	45.50
20	ATOM	809	CG	PRO	100	29.750	74.757	7.870	1.00	45.80
	ATOM	810	C	PRO	100	26.727	76.553	6.796	1.00	46.78
	ATOM	811	O	PRO	100	26.292	75.737	5.994	1.00	52.13
	ATOM	812	N	ASN	101	26.249	77.781	6.901	1.00	42.71
	ATOM	813	CA	ASN	101	25.184	78.269	6.046	1.00	41.37
25	ATOM	814	CB	ASN	101	23.846	77.677	6.470	1.00	43.24
	ATOM	815	CG	ASN	101	23.600	76.304	5.878	1.00	45.93
	ATOM	816	OD1	ASN	101	23.387	76.163	4.672	1.00	45.33
	ATOM	817	ND2	ASN	101	23.634	75.277	6.726	1.00	48.11
	ATOM	818	C	ASN	101	25.145	79.780	6.160	1.00	42.66
30	ATOM	819	O	ASN	101	24.348	80.336	6.918	1.00	44.79
	ATOM	820	N	ASN	102	26.012	80.460	5.415	1.00	40.46
	ATOM	821	CA	ASN	102	26.014	81.902	5.496	1.00	37.90
	ATOM	822	CB	ASN	102	27.375	82.408	5.949	1.00	34.01
	ATOM	823	CG	ASN	102	27.493	82.448	7.460	1.00	32.44
35	ATOM	824	OD1	ASN	102	26.549	82.824	8.151	1.00	29.07
	ATOM	825	ND2	ASN	102	28.650	82.067	7.979	1.00	33.79
	ATOM	826	C	ASN	102	25.531	82.710	4.308	1.00	39.20
	ATOM	827	O	ASN	102	24.525	83.401	4.416	1.00	46.07
	ATOM	828	N	SER	103	26.201	82.649	3.174	1.00	35.42
40	ATOM	829	CA	SER	103	25.742	83.472	2.041	1.00	40.28
	ATOM	830	CB	SER	103	24.244	83.238	1.728	1.00	40.68
	ATOM	831	OG	SER	103	23.407	84.278	2.220	1.00	31.76
	ATOM	832	C	SER	103	25.989	84.973	2.323	1.00	35.24
	ATOM	833	O	SER	103	25.522	85.536	3.311	1.00	24.65
45	ATOM	834	N	CYS	104	26.763	85.595	1.443	1.00	32.88
	ATOM	835	CA	CYS	104	27.094	86.991	1.568	1.00	31.77
	ATOM	836	C	CYS	104	26.719	87.689	0.248	1.00	30.58
	ATOM	837	O	CYS	104	26.982	87.176	-0.836	1.00	29.23
	ATOM	838	CB	CYS	104	28.605	87.168	1.886	1.00	28.43
50	ATOM	839	SG	CYS	104	29.026	88.901	1.644	1.00	46.07
	ATOM	840	N	TYR	105	26.070	88.847	0.350	1.00	30.34
	ATOM	841	CA	TYR	105	25.685	89.642	-0.815	1.00	26.12
	ATOM	842	CB	TYR	105	24.211	90.035	-0.736	1.00	27.03
	ATOM	843	CG	TYR	105	23.755	91.022	-1.799	1.00	25.61
55	ATOM	844	CD1	TYR	105	23.110	90.591	-2.963	1.00	24.90
	ATOM	845	CE1	TYR	105	22.696	91.492	-3.934	1.00	26.08
	ATOM	846	CD2	TYR	105	23.971	92.388	-1.645	1.00	25.41
	ATOM	847	CE2	TYR	105	23.562	93.294	-2.615	1.00	28.91
	ATOM	848	CZ	TYR	105	22.925	92.840	-3.755	1.00	27.58
60	ATOM	849	OH	TYR	105	22.509	93.744	-4.708	1.00	29.89
	ATOM	850	C	TYR	105	26.535	90.909	-0.829	1.00	29.20
	ATOM	851	O	TYR	105	26.851	91.467	0.217	1.00	33.10
	ATOM	852	N	SER	106	26.910	91.368	-2.012	1.00	30.48
	ATOM	853	CA	SER	106	27.711	92.577	-2.112	1.00	29.22
65	ATOM	854	CB	SER	106	29.190	92.246	-1.896	1.00	28.30
	ATOM	855	OG	SER	106	29.996	93.407	-1.935	1.00	28.62

	ATOM	856	C	SER	106	27.495	93.171	-3.493	1.00	29.94
	ATOM	857	O	SER	106	27.354	92.434	-4.461	1.00	29.72
	ATOM	858	N	ALA	107	27.445	94.498	-3.579	1.00	27.64
	ATOM	859	CA	ALA	107	27.244	95.173	-4.854	1.00	24.33
5	ATOM	860	CB	ALA	107	25.762	95.249	-5.178	1.00	21.30
	ATOM	861	C	ALA	107	27.846	96.569	-4.856	1.00	25.56
	ATOM	862	O	ALA	107	28.184	97.117	-3.816	1.00	28.03
	ATOM	863	N	GLY	108	27.976	97.142	-6.042	1.00	26.07
	ATOM	864	CA	GLY	108	28.525	98.477	-6.161	1.00	25.49
10	ATOM	865	C	GLY	108	28.481	98.935	-7.600	1.00	29.03
	ATOM	866	O	GLY	108	28.090	98.176	-8.476	1.00	32.20
	ATOM	867	N	ILE	109	28.880	100.174	-7.850	1.00	29.34
	ATOM	868	CA	ILE	109	28.879	100.719	-9.200	1.00	27.71
	ATOM	869	CB	ILE	109	28.227	102.111	-9.230	1.00	27.48
15	ATOM	870	CG2	ILE	109	28.253	102.673	-10.635	1.00	26.46
	ATOM	871	CG1	ILE	109	26.789	102.009	-8.724	1.00	26.16
	ATOM	872	CD1	ILE	109	26.106	103.336	-8.552	1.00	29.07
	ATOM	873	C	ILE	109	30.303	100.827	-9.709	1.00	28.66
	ATOM	874	O	ILE	109	31.218	101.148	-8.953	1.00	30.59
20	ATOM	875	N	ALA	110	30.496	100.554	-10.992	1.00	25.60
	ATOM	876	CA	ALA	110	31.819	100.630	-11.599	1.00	26.36
	ATOM	877	CB	ALA	110	32.508	99.288	-11.510	1.00	20.26
	ATOM	878	C	ALA	110	31.697	101.045	-13.054	1.00	29.61
	ATOM	879	O	ALA	110	30.669	100.832	-13.678	1.00	27.61
25	ATOM	880	N	LYS	111	32.738	101.661	-13.596	1.00	33.77
	ATOM	881	CA	LYS	111	32.692	102.046	-14.987	1.00	35.09
	ATOM	882	CB	LYS	111	33.396	103.376	-15.225	1.00	39.38
	ATOM	883	CG	LYS	111	33.482	103.688	-16.714	1.00	49.74
	ATOM	884	CD	LYS	111	33.371	105.156	-17.020	1.00	53.95
30	ATOM	885	CE	LYS	111	33.290	105.370	-18.524	1.00	54.28
	ATOM	886	NZ	LYS	111	33.206	106.834	-18.852	1.00	60.85
	ATOM	887	C	LYS	111	33.360	100.946	-15.804	1.00	35.28
	ATOM	888	O	LYS	111	34.489	100.556	-15.529	1.00	36.58
	ATOM	889	N	LEU	112	32.643	100.445	-16.801	1.00	32.32
35	ATOM	890	CA	LEU	112	33.142	99.381	-17.650	1.00	29.67
	ATOM	891	CB	LEU	112	32.237	98.157	-17.530	1.00	27.04
	ATOM	892	CG	LEU	112	31.909	97.695	-16.111	1.00	24.38
	ATOM	893	CD1	LEU	112	30.899	96.567	-16.158	1.00	27.56
	ATOM	894	CD2	LEU	112	33.167	97.258	-15.412	1.00	19.52
40	ATOM	895	C	LEU	112	33.184	99.840	-19.099	1.00	32.58
	ATOM	896	O	LEU	112	32.582	100.853	-19.455	1.00	31.51
	ATOM	897	N	GLU	113	33.891	99.089	-19.935	1.00	35.75
	ATOM	898	CA	GLU	113	34.008	99.426	-21.342	1.00	40.08
	ATOM	899	CB	GLU	113	35.402	99.932	-21.664	1.00	44.23
45	ATOM	900	CG	GLU	113	35.955	100.965	-20.727	1.00	50.12
	ATOM	901	CD	GLU	113	37.059	101.752	-21.394	1.00	58.08
	ATOM	902	OE1	GLU	113	37.856	101.134	-22.160	1.00	61.13
	ATOM	903	OE2	GLU	113	37.126	102.985	-21.152	1.00	63.64
	ATOM	904	C	GLU	113	33.756	98.234	-22.231	1.00	41.63
50	ATOM	905	O	GLU	113	33.853	97.095	-21.795	1.00	48.29
	ATOM	906	N	GLU	114	33.446	98.516	-23.492	1.00	40.97
	ATOM	907	CA	GLU	114	33.217	97.486	-24.496	1.00	36.68
	ATOM	908	CB	GLU	114	33.247	98.091	-25.883	1.00	41.19
	ATOM	909	CG	GLU	114	31.935	98.443	-26.487	1.00	46.58
55	ATOM	910	CD	GLU	114	32.027	98.414	-27.997	1.00	51.12
	ATOM	911	OE1	GLU	114	32.222	97.303	-28.550	1.00	51.88
	ATOM	912	OE2	GLU	114	31.922	99.495	-28.625	1.00	57.08
	ATOM	913	C	GLU	114	34.358	96.492	-24.442	1.00	35.51
	ATOM	914	O	GLU	114	35.512	96.872	-24.600	1.00	38.92
60	ATOM	915	N	GLY	115	34.046	95.222	-24.238	1.00	32.99
	ATOM	916	CA	GLY	115	35.104	94.237	-24.207	1.00	31.79
	ATOM	917	C	GLY	115	35.499	93.803	-22.821	1.00	34.39
	ATOM	918	O	GLY	115	36.149	92.778	-22.668	1.00	39.20
	ATOM	919	N	ASP	116	35.138	94.587	-21.814	1.00	35.66
65	ATOM	920	CA	ASP	116	35.455	94.219	-20.445	1.00	38.09
	ATOM	921	CB	ASP	116	35.094	95.343	-19.467	1.00	36.00

	ATOM	922	CG	ASP	116	36.095	96.477	-19.471	1.00	35.09
	ATOM	923	OD1	ASP	116	37.215	96.300	-19.993	1.00	33.81
	ATOM	924	OD2	ASP	116	35.765	97.549	-18.931	1.00	35.24
	ATOM	925	C	ASP	116	34.630	92.988	-20.103	1.00	39.38
5	ATOM	926	O	ASP	116	33.559	92.767	-20.675	1.00	37.91
	ATOM	927	N	GLU	117	35.132	92.176	-19.181	1.00	41.54
	ATOM	928	CA	GLU	117	34.398	90.992	-18.754	1.00	37.92
	ATOM	929	CB	GLU	117	35.114	89.719	-19.189	1.00	38.90
	ATOM	930	CG	GLU	117	35.328	89.614	-20.677	1.00	48.15
10	ATOM	931	CD	GLU	117	36.004	88.316	-21.065	1.00	52.62
	ATOM	932	OE1	GLU	117	36.903	87.854	-20.309	1.00	55.49
	ATOM	933	OE2	GLU	117	35.641	87.765	-22.133	1.00	57.10
	ATOM	934	C	GLU	117	34.301	91.021	-17.241	1.00	34.02
	ATOM	935	O	GLU	117	35.223	91.477	-16.567	1.00	33.31
15	ATOM	936	N	LEU	118	33.174	90.563	-16.714	1.00	30.68
	ATOM	937	CA	LEU	118	32.971	90.506	-15.276	1.00	27.14
	ATOM	938	CB	LEU	118	31.649	91.172	-14.893	1.00	26.49
	ATOM	939	CG	LEU	118	31.499	92.672	-15.120	1.00	26.21
	ATOM	940	CD1	LEU	118	30.095	93.094	-14.755	1.00	23.05
20	ATOM	941	CD2	LEU	118	32.515	93.425	-14.288	1.00	25.38
	ATOM	942	C	LEU	118	32.926	89.042	-14.858	1.00	28.33
	ATOM	943	O	LEU	118	32.433	88.195	-15.604	1.00	31.46
	ATOM	944	N	GLN	119	33.448	88.735	-13.675	1.00	28.96
	ATOM	945	CA	GLN	119	33.406	87.364	-13.182	1.00	29.13
25	ATOM	946	CB	GLN	119	34.623	86.590	-13.667	1.00	28.23
	ATOM	947	CG	GLN	119	35.913	87.019	-13.038	1.00	36.38
	ATOM	948	CD	GLN	119	37.099	86.272	-13.599	1.00	36.57
	ATOM	949	OE1	GLN	119	38.180	86.283	-13.011	1.00	40.70
	ATOM	950	NE2	GLN	119	36.910	85.625	-14.743	1.00	36.80
30	ATOM	951	C	GLN	119	33.316	87.307	-11.660	1.00	29.43
	ATOM	952	O	GLN	119	33.744	88.227	-10.967	1.00	28.80
	ATOM	953	N	LEU	120	32.735	86.222	-11.161	1.00	32.26
	ATOM	954	CA	LEU	120	32.553	85.984	-9.732	1.00	29.96
	ATOM	955	CB	LEU	120	31.102	85.557	-9.466	1.00	31.17
35	ATOM	956	CG	LEU	120	30.446	85.475	-8.081	1.00	30.26
	ATOM	957	CD1	LEU	120	31.335	84.758	-7.096	1.00	27.10
	ATOM	958	CD2	LEU	120	30.138	86.868	-7.609	1.00	30.96
	ATOM	959	C	LEU	120	33.510	84.853	-9.349	1.00	30.51
	ATOM	960	O	LEU	120	33.349	83.714	-9.790	1.00	32.88
40	ATOM	961	N	ALA	121	34.501	85.169	-8.524	1.00	27.40
	ATOM	962	CA	ALA	121	35.487	84.180	-8.109	1.00	25.55
	ATOM	963	CB	ALA	121	36.859	84.593	-8.616	1.00	19.41
	ATOM	964	C	ALA	121	35.547	83.944	-6.599	1.00	29.89
	ATOM	965	O	ALA	121	35.472	84.881	-5.807	1.00	35.88
45	ATOM	966	N	ILE	122	35.675	82.678	-6.214	1.00	30.33
	ATOM	967	CA	ILE	122	35.783	82.305	-4.810	1.00	30.27
	ATOM	968	CB	ILE	122	34.849	81.143	-4.460	1.00	29.27
	ATOM	969	CG2	ILE	122	34.933	80.859	-2.978	1.00	30.38
	ATOM	970	CG1	ILE	122	33.411	81.495	-4.851	1.00	26.03
50	ATOM	971	CD1	ILE	122	32.394	80.402	-4.580	1.00	29.21
	ATOM	972	C	ILE	122	37.230	81.867	-4.606	1.00	33.20
	ATOM	973	O	ILE	122	37.666	80.868	-5.174	1.00	33.30
	ATOM	974	N	PRO	123	37.995	82.617	-3.794	1.00	34.70
	ATOM	975	CD	PRO	123	37.565	83.842	-3.113	1.00	34.40
55	ATOM	976	CA	PRO	123	39.408	82.343	-3.496	1.00	36.49
	ATOM	977	CB	PRO	123	39.887	83.636	-2.828	1.00	30.22
	ATOM	978	CG	PRO	123	38.822	84.645	-3.149	1.00	36.72
	ATOM	979	C	PRO	123	39.636	81.141	-2.581	1.00	39.64
	ATOM	980	O	PRO	123	40.258	81.277	-1.523	1.00	40.48
60	ATOM	981	N	ARG	124	39.138	79.976	-2.986	1.00	44.58
	ATOM	982	CA	ARG	124	39.294	78.763	-2.196	1.00	48.04
	ATOM	983	CB	ARG	124	38.141	78.622	-1.209	1.00	52.91
	ATOM	984	CG	ARG	124	37.952	79.829	-0.310	1.00	60.50
	ATOM	985	CD	ARG	124	38.378	79.466	1.096	1.00	72.57
65	ATOM	986	NE	ARG	124	38.378	80.636	1.993	1.00	84.90
	ATOM	987	CZ	ARG	124	39.426	81.439	2.217	1.00	85.44

	ATOM	988	NH1	ARG	124	40.588	81.211	1.604	1.00	87.69
	ATOM	989	NH2	ARG	124	39.307	82.477	3.055	1.00	82.78
	ATOM	990	C	ARG	124	39.374	77.524	-3.073	1.00	48.39
	ATOM	991	O	ARG	124	38.758	77.469	-4.138	1.00	45.26
5	ATOM	992	N	GLU	125	40.134	76.539	-2.601	1.00	52.84
	ATOM	993	CA	GLU	125	40.357	75.281	-3.309	1.00	55.02
	ATOM	994	CB	GLU	125	41.128	74.314	-2.406	1.00	63.62
	ATOM	995	CG	GLU	125	40.869	74.517	-0.911	1.00	70.62
	ATOM	996	CD	GLU	125	41.141	75.946	-0.457	1.00	74.76
10	ATOM	997	OE1	GLU	125	42.324	76.379	-0.514	1.00	75.75
	ATOM	998	OE2	GLU	125	40.173	76.632	-0.047	1.00	80.71
	ATOM	999	C	GLU	125	39.102	74.603	-3.841	1.00	53.44
	ATOM	1000	O	GLU	125	38.893	74.554	-5.053	1.00	57.50
	ATOM	1001	N	ASN	126	38.288	74.039	-2.959	1.00	47.18
15	ATOM	1002	CA	ASN	126	37.056	73.409	-3.416	1.00	47.59
	ATOM	1003	CB	ASN	126	37.131	71.887	-3.332	1.00	51.03
	ATOM	1004	CG	ASN	126	37.398	71.246	-4.686	1.00	57.07
	ATOM	1005	OD1	ASN	126	38.527	71.270	-5.199	1.00	61.15
	ATOM	1006	ND2	ASN	126	36.348	70.684	-5.286	1.00	56.63
20	ATOM	1007	C	ASN	126	35.933	73.924	-2.558	1.00	47.06
	ATOM	1008	O	ASN	126	35.365	73.206	-1.731	1.00	45.61
	ATOM	1009	N	ALA	127	35.632	75.199	-2.766	1.00	43.50
	ATOM	1010	CA	ALA	127	34.604	75.884	-2.016	1.00	38.26
	ATOM	1011	CB	ALA	127	34.309	77.216	-2.666	1.00	37.97
25	ATOM	1012	C	ALA	127	33.330	75.070	-1.902	1.00	37.16
	ATOM	1013	O	ALA	127	32.796	74.584	-2.897	1.00	33.14
	ATOM	1014	N	GLN	128	32.862	74.904	-0.670	1.00	37.87
	ATOM	1015	CA	GLN	128	31.614	74.195	-0.430	1.00	37.19
	ATOM	1016	CB	GLN	128	31.636	73.564	0.958	1.00	40.23
30	ATOM	1017	CG	GLN	128	32.633	72.433	1.051	1.00	38.14
	ATOM	1018	CD	GLN	128	32.391	71.403	-0.027	1.00	39.51
	ATOM	1019	OE1	GLN	128	31.376	70.698	-0.017	1.00	40.36
	ATOM	1020	NE2	GLN	128	33.313	71.322	-0.986	1.00	41.53
	ATOM	1021	C	GLN	128	30.534	75.268	-0.549	1.00	35.55
35	ATOM	1022	O	GLN	128	30.291	76.055	0.370	1.00	35.57
	ATOM	1023	N	ILE	129	29.903	75.284	-1.714	1.00	34.64
	ATOM	1024	CA	ILE	129	28.903	76.280	-2.072	1.00	29.60
	ATOM	1025	CB	ILE	129	29.487	77.124	-3.263	1.00	31.03
	ATOM	1026	CG2	ILE	129	28.494	77.301	-4.382	1.00	29.76
40	ATOM	1027	CG1	ILE	129	30.004	78.444	-2.735	1.00	26.37
	ATOM	1028	CD1	ILE	129	31.093	78.277	-1.744	1.00	34.70
	ATOM	1029	C	ILE	129	27.559	75.668	-2.452	1.00	29.55
	ATOM	1030	O	ILE	129	27.492	74.524	-2.877	1.00	33.42
	ATOM	1031	N	SER	130	26.482	76.428	-2.288	1.00	27.67
45	ATOM	1032	CA	SER	130	25.161	75.947	-2.681	1.00	25.00
	ATOM	1033	CB	SER	130	24.078	76.516	-1.780	1.00	22.09
	ATOM	1034	OG	SER	130	22.797	76.227	-2.322	1.00	20.89
	ATOM	1035	C	SER	130	24.904	76.414	-4.104	1.00	26.03
	ATOM	1036	O	SER	130	25.180	77.556	-4.429	1.00	30.77
50	ATOM	1037	N	LEU	131	24.370	75.547	-4.951	1.00	24.49
	ATOM	1038	CA	LEU	131	24.113	75.922	-6.332	1.00	28.07
	ATOM	1039	CB	LEU	131	24.711	74.884	-7.283	1.00	28.87
	ATOM	1040	CG	LEU	131	26.165	75.101	-7.700	1.00	30.76
	ATOM	1041	CD1	LEU	131	27.031	75.401	-6.512	1.00	28.98
55	ATOM	1042	CD2	LEU	131	26.654	73.866	-8.410	1.00	33.37
	ATOM	1043	C	LEU	131	22.647	76.136	-6.672	1.00	33.66
	ATOM	1044	O	LEU	131	22.229	75.902	-7.806	1.00	37.67
	ATOM	1045	N	ASP	132	21.868	76.578	-5.691	1.00	34.97
	ATOM	1046	CA	ASP	132	20.452	76.850	-5.915	1.00	34.31
60	ATOM	1047	CB	ASP	132	19.667	76.772	-4.606	1.00	42.68
	ATOM	1048	CG	ASP	132	19.425	75.341	-4.153	1.00	47.98
	ATOM	1049	OD1	ASP	132	18.979	75.152	-2.994	1.00	50.41
	ATOM	1050	OD2	ASP	132	19.671	74.412	-4.963	1.00	46.42
	ATOM	1051	C	ASP	132	20.322	78.239	-6.517	1.00	34.32
65	ATOM	1052	O	ASP	132	20.933	79.191	-6.043	1.00	32.87
	ATOM	1053	N	GLY	133	19.521	78.341	-7.569	1.00	34.53

	ATOM	1054	CA	GLY	133	19.340	79.607	-8.250	1.00	36.93
	ATOM	1055	C	GLY	133	18.911	80.793	-7.410	1.00	35.06
	ATOM	1056	O	GLY	133	19.127	81.940	-7.802	1.00	40.44
	ATOM	1057	N	ASP	134	18.312	80.541	-6.256	1.00	31.08
5	ATOM	1058	CA	ASP	134	17.857	81.638	-5.426	1.00	27.63
	ATOM	1059	CB	ASP	134	16.521	81.280	-4.768	1.00	28.43
	ATOM	1060	CG	ASP	134	16.605	80.057	-3.882	1.00	31.78
	ATOM	1061	OD1	ASP	134	17.382	79.137	-4.201	1.00	35.68
	ATOM	1062	OD2	ASP	134	15.876	80.005	-2.870	1.00	32.53
10	ATOM	1063	C	ASP	134	18.869	82.102	-4.394	1.00	25.08
	ATOM	1064	O	ASP	134	18.797	83.218	-3.930	1.00	19.71
	ATOM	1065	N	VAL	135	19.842	81.271	-4.057	1.00	26.84
	ATOM	1066	CA	VAL	135	20.824	81.690	-3.068	1.00	23.72
	ATOM	1067	CB	VAL	135	21.100	80.580	-2.051	1.00	25.13
15	ATOM	1068	CG1	VAL	135	19.889	80.405	-1.175	1.00	22.76
	ATOM	1069	CG2	VAL	135	21.440	79.287	-2.761	1.00	22.12
	ATOM	1070	C	VAL	135	22.144	82.205	-3.631	1.00	26.54
	ATOM	1071	O	VAL	135	22.826	82.977	-2.970	1.00	29.26
	ATOM	1072	N	THR	136	22.523	81.785	-4.836	1.00	26.39
20	ATOM	1073	CA	THR	136	23.758	82.304	-5.422	1.00	28.40
	ATOM	1074	CB	THR	136	24.936	81.277	-5.322	1.00	28.21
	ATOM	1075	OG1	THR	136	25.154	80.651	-6.584	1.00	37.13
	ATOM	1076	CG2	THR	136	24.640	80.223	-4.285	1.00	26.90
	ATOM	1077	C	THR	136	23.520	82.760	-6.869	1.00	31.60
25	ATOM	1078	O	THR	136	23.163	81.967	-7.745	1.00	27.29
	ATOM	1079	N	PHE	137	23.702	84.062	-7.094	1.00	31.45
	ATOM	1080	CA	PHE	137	23.473	84.670	-8.400	1.00	26.64
	ATOM	1081	CB	PHE	137	22.040	85.179	-8.455	1.00	26.84
	ATOM	1082	CG	PHE	137	21.598	85.878	-7.203	1.00	24.32
30	ATOM	1083	CD1	PHE	137	21.847	87.234	-7.019	1.00	27.77
	ATOM	1084	CD2	PHE	137	20.923	85.187	-6.210	1.00	23.70
	ATOM	1085	CE1	PHE	137	21.431	87.888	-5.871	1.00	27.32
	ATOM	1086	CE2	PHE	137	20.505	85.834	-5.057	1.00	26.31
	ATOM	1087	CZ	PHE	137	20.759	87.188	-4.891	1.00	29.43
35	ATOM	1088	C	PHE	137	24.469	85.785	-8.710	1.00	28.06
	ATOM	1089	O	PHE	137	25.157	86.263	-7.820	1.00	25.04
	ATOM	1090	N	PHE	138	24.532	86.198	-9.975	1.00	29.81
	ATOM	1091	CA	PHE	138	25.491	87.211	-10.411	1.00	29.93
	ATOM	1092	CB	PHE	138	26.579	86.503	-11.215	1.00	27.47
40	ATOM	1093	CG	PHE	138	27.783	87.337	-11.491	1.00	28.49
	ATOM	1094	CD1	PHE	138	28.109	88.418	-10.679	1.00	27.65
	ATOM	1095	CD2	PHE	138	28.607	87.034	-12.572	1.00	31.06
	ATOM	1096	CE1	PHE	138	29.235	89.186	-10.944	1.00	30.90
	ATOM	1097	CE2	PHE	138	29.737	87.793	-12.848	1.00	27.65
45	ATOM	1098	CZ	PHE	138	30.052	88.872	-12.034	1.00	27.41
	ATOM	1099	C	PHE	138	24.859	88.379	-11.194	1.00	30.91
	ATOM	1100	O	PHE	138	24.140	88.185	-12.167	1.00	24.94
	ATOM	1101	N	GLY	139	25.191	89.591	-10.739	1.00	36.94
	ATOM	1102	CA	GLY	139	24.666	90.865	-11.228	1.00	36.08
50	ATOM	1103	C	GLY	139	24.752	91.499	-12.594	1.00	36.48
	ATOM	1104	O	GLY	139	24.453	90.860	-13.586	1.00	43.61
	ATOM	1105	N	ALA	140	25.091	92.789	-12.613	1.00	36.60
	ATOM	1106	CA	ALA	140	25.227	93.615	-13.828	1.00	34.94
	ATOM	1107	CB	ALA	140	25.943	92.836	-14.917	1.00	36.94
55	ATOM	1108	C	ALA	140	23.958	94.263	-14.415	1.00	32.53
	ATOM	1109	O	ALA	140	23.142	93.611	-15.056	1.00	27.96
	ATOM	1110	N	LEU	141	23.832	95.571	-14.204	1.00	33.23
	ATOM	1111	CA	LEU	141	22.705	96.375	-14.697	1.00	36.11
	ATOM	1112	CB	LEU	141	21.702	96.606	-13.567	1.00	37.53
60	ATOM	1113	CG	LEU	141	20.468	97.490	-13.771	1.00	38.87
	ATOM	1114	CD1	LEU	141	19.547	97.317	-12.588	1.00	40.67
	ATOM	1115	CD2	LEU	141	20.848	98.952	-13.904	1.00	37.53
	ATOM	1116	C	LEU	141	23.251	97.720	-15.186	1.00	38.89
	ATOM	1117	O	LEU	141	23.984	98.392	-14.465	1.00	39.25
65	ATOM	1118	N	LYS	142	22.897	98.130	-16.400	1.00	42.86
	ATOM	1119	CA	LYS	142	23.413	99.397	-16.907	1.00	42.96

	ATOM	1120	CB	LYS	142	23.493	99.366	-18.430	1.00	41.80
	ATOM	1121	CG	LYS	142	23.996	100.670	-19.002	1.00	46.88
	ATOM	1122	CD	LYS	142	24.528	100.524	-20.412	1.00	49.58
	ATOM	1123	CE	LYS	142	25.016	101.873	-20.913	1.00	51.07
5	ATOM	1124	NZ	LYS	142	25.629	101.791	-22.261	1.00	56.20
	ATOM	1125	C	LYS	142	22.640	100.633	-16.462	1.00	43.41
	ATOM	1126	O	LYS	142	21.420	100.694	-16.588	1.00	44.98
	ATOM	1127	N	LEU	143	23.363	101.622	-15.945	1.00	42.81
	ATOM	1128	CA	LEU	143	22.751	102.866	-15.484	1.00	40.65
10	ATOM	1129	CB	LEU	143	23.630	103.529	-14.424	1.00	36.56
	ATOM	1130	CG	LEU	143	23.982	102.731	-13.168	1.00	36.17
	ATOM	1131	CD1	LEU	143	24.906	103.559	-12.295	1.00	37.95
	ATOM	1132	CD2	LEU	143	22.721	102.367	-12.414	1.00	27.38
	ATOM	1133	C	LEU	143	22.580	103.839	-16.645	1.00	43.14
15	ATOM	1134	O	LEU	143	23.378	103.838	-17.589	1.00	46.28
	ATOM	1135	N	LEU	144	21.547	104.668	-16.582	1.00	42.63
	ATOM	1136	CA	LEU	144	21.316	105.642	-17.638	1.00	44.11
	ATOM	1137	CB	LEU	144	19.869	106.124	-17.617	1.00	44.88
	ATOM	1138	CG	LEU	144	18.839	105.044	-17.942	1.00	47.02
20	ATOM	1139	CD1	LEU	144	17.448	105.599	-17.795	1.00	47.71
	ATOM	1140	CD2	LEU	144	19.058	104.534	-19.354	1.00	47.87
	ATOM	1141	C	LEU	144	22.240	106.830	-17.452	1.00	45.66
	ATOM	1142	O	LEU	144	22.732	107.025	-16.325	1.00	46.37
	ATOM	1143	OXT	LEU	144	22.450	107.560	-18.439	1.00	50.66
25	END					134.002	98.540	-12.573	0.00	0.00

TABLE 7

111111									
	ATOM	1	CB	VAL	1	19.384	112.619	-19.765	1.00 69.28
5	ATOM	2	CG1	VAL	1	19.749	111.864	-21.062	1.00 68.26
	ATOM	3	CG2	VAL	1	18.590	113.894	-20.070	1.00 71.75
	ATOM	4	C	VAL	1	19.532	110.644	-18.221	1.00 64.86
	ATOM	5	O	VAL	1	19.522	109.473	-18.629	1.00 64.02
	ATOM	6	N	VAL	1	17.901	112.472	-17.726	1.00 65.10
10	ATOM	7	CA	VAL	1	18.566	111.689	-18.812	1.00 65.98
	ATOM	8	N	THR	2	20.369	111.071	-17.272	1.00 61.42
	ATOM	9	CA	THR	2	21.326	110.165	-16.638	1.00 56.64
	ATOM	10	CB	THR	2	22.780	110.623	-16.849	1.00 56.77
	ATOM	11	OG1	THR	2	22.919	111.967	-16.380	1.00 57.45
15	ATOM	12	CG2	THR	2	23.168	110.553	-18.321	1.00 55.68
	ATOM	13	C	THR	2	21.103	110.074	-15.141	1.00 53.08
	ATOM	14	O	THR	2	20.351	110.852	-14.563	1.00 52.00
	ATOM	15	N	GLN	3	21.783	109.123	-14.513	1.00 50.34
	ATOM	16	CA	GLN	3	21.666	108.924	-13.074	1.00 45.97
20	ATOM	17	CB	GLN	3	21.329	107.468	-12.774	1.00 46.09
	ATOM	18	CG	GLN	3	20.188	106.924	-13.593	1.00 49.60
	ATOM	19	CD	GLN	3	19.839	105.510	-13.205	1.00 50.43
	ATOM	20	OE1	GLN	3	19.350	105.270	-12.106	1.00 50.94
	ATOM	21	NE2	GLN	3	20.096	104.561	-14.103	1.00 49.99
25	ATOM	22	C	GLN	3	22.958	109.292	-12.369	1.00 42.41
	ATOM	23	O	GLN	3	23.966	108.601	-12.488	1.00 37.34
	ATOM	24	N	ASP	4	22.925	110.390	-11.632	1.00 40.32
	ATOM	25	CA	ASP	4	24.102	110.837	-10.909	1.00 39.37
	ATOM	26	CB	ASP	4	23.863	112.231	-10.323	1.00 44.56
30	ATOM	27	CG	ASP	4	23.694	113.290	-11.392	1.00 45.76
	ATOM	28	OD1	ASP	4	23.399	112.915	-12.550	1.00 48.58
	ATOM	29	OD2	ASP	4	23.847	114.489	-11.075	1.00 45.55
	ATOM	30	C	ASP	4	24.396	109.861	-9.791	1.00 36.30
	ATOM	31	O	ASP	4	23.487	109.270	-9.214	1.00 36.29
35	ATOM	32	N	CYS	5	25.674	109.685	-9.496	1.00 35.11
	ATOM	33	CA	CYS	5	26.086	108.803	-8.419	1.00 35.61
	ATOM	34	CB	CYS	5	25.986	107.334	-8.839	1.00 34.96
	ATOM	35	SG	CYS	5	26.567	106.970	-10.483	1.00 35.34
	ATOM	36	C	CYS	5	27.497	109.147	-7.991	1.00 31.80
40	ATOM	37	O	CYS	5	28.288	109.654	-8.777	1.00 31.85
	ATOM	38	N	LEU	6	27.790	108.903	-6.724	1.00 27.72
	ATOM	39	CA	LEU	6	29.102	109.170	-6.172	1.00 30.80
	ATOM	40	CB	LEU	6	29.087	110.474	-5.383	1.00 29.25
	ATOM	41	CG	LEU	6	30.372	110.883	-4.661	1.00 29.47
45	ATOM	42	CD1	LEU	6	30.398	112.377	-4.479	1.00 30.65
	ATOM	43	CD2	LEU	6	30.442	110.206	-3.325	1.00 35.27
	ATOM	44	C	LEU	6	29.451	108.006	-5.266	1.00 31.83
	ATOM	45	O	LEU	6	28.668	107.644	-4.396	1.00 33.45
	ATOM	46	N	GLN	7	30.619	107.410	-5.475	1.00 32.23
50	ATOM	47	CA	GLN	7	31.032	106.280	-4.671	1.00 30.26
	ATOM	48	CB	GLN	7	31.221	105.059	-5.562	1.00 29.56
	ATOM	49	CG	GLN	7	31.364	103.746	-4.824	1.00 25.09
	ATOM	50	CD	GLN	7	31.288	102.562	-5.767	1.00 26.50
	ATOM	51	OE1	GLN	7	32.200	102.314	-6.556	1.00 24.40
55	ATOM	52	NE2	GLN	7	30.188	101.834	-5.702	1.00 27.03
	ATOM	53	C	GLN	7	32.314	106.601	-3.934	1.00 32.11
	ATOM	54	O	GLN	7	33.206	107.247	-4.469	1.00 34.24
	ATOM	55	N	LEU	8	32.382	106.153	-2.689	1.00 33.73
	ATOM	56	CA	LEU	8	33.542	106.371	-1.840	1.00 35.41
60	ATOM	57	CB	LEU	8	33.122	107.105	-0.555	1.00 32.44
	ATOM	58	CG	LEU	8	33.191	108.637	-0.468	1.00 27.69
	ATOM	59	CD1	LEU	8	33.251	109.272	-1.833	1.00 26.43
	ATOM	60	CD2	LEU	8	32.002	109.122	0.315	1.00 18.64
	ATOM	61	C	LEU	8	34.197	105.041	-1.498	1.00 36.87
65	ATOM	62	O	LEU	8	33.533	104.014	-1.369	1.00 36.05
	ATOM	63	N	ILE	9	35.512	105.079	-1.344	1.00 39.25

	ATOM	64	CA	ILE	9	36.303	103.899	-1.038	1.00	35.57
	ATOM	65	CB	ILE	9	37.196	103.570	-2.261	1.00	34.28
	ATOM	66	CG2	ILE	9	38.616	103.270	-1.852	1.00	38.20
	ATOM	67	CG1	ILE	9	36.599	102.411	-3.025	1.00	35.22
5	ATOM	68	CD1	ILE	9	37.456	102.017	-4.180	1.00	43.63
	ATOM	69	C	ILE	9	37.146	104.158	0.214	1.00	35.28
	ATOM	70	O	ILE	9	37.570	105.284	0.458	1.00	33.75
	ATOM	71	N	ALA	10	37.376	103.126	1.017	1.00	35.78
10	ATOM	72	CA	ALA	10	38.179	103.298	2.218	1.00	35.28
	ATOM	73	CB	ALA	10	38.203	102.018	3.021	1.00	28.47
	ATOM	74	C	ALA	10	39.600	103.709	1.855	1.00	36.49
	ATOM	75	O	ALA	10	40.184	103.200	0.903	1.00	37.41
	ATOM	76	N	ASP	11	40.152	104.639	2.621	1.00	39.44
	ATOM	77	CA	ASP	11	41.510	105.116	2.389	1.00	40.77
15	ATOM	78	CB	ASP	11	41.544	106.644	2.445	1.00	38.90
	ATOM	79	CG	ASP	11	42.946	107.193	2.342	1.00	40.34
	ATOM	80	OD1	ASP	11	43.820	106.508	1.758	1.00	39.50
	ATOM	81	OD2	ASP	11	43.167	108.314	2.837	1.00	37.38
	ATOM	82	C	ASP	11	42.456	104.528	3.430	1.00	42.25
20	ATOM	83	O	ASP	11	42.618	105.069	4.525	1.00	40.54
	ATOM	84	N	SER	12	43.083	103.416	3.068	1.00	42.92
	ATOM	85	CA	SER	12	43.994	102.699	3.958	1.00	43.84
	ATOM	86	CB	SER	12	44.433	101.397	3.298	1.00	42.13
	ATOM	87	OG	SER	12	45.059	101.659	2.055	1.00	43.31
25	ATOM	88	C	SER	12	45.230	103.475	4.381	1.00	46.10
	ATOM	89	O	SER	12	46.016	102.998	5.200	1.00	43.90
	ATOM	90	N	GLU	13	45.400	104.672	3.832	1.00	46.72
	ATOM	91	CA	GLU	13	46.560	105.476	4.165	1.00	44.93
	ATOM	92	CB	GLU	13	47.172	106.046	2.898	1.00	46.71
30	ATOM	93	CG	GLU	13	47.759	104.967	2.030	1.00	58.25
	ATOM	94	CD	GLU	13	48.850	105.494	1.121	1.00	66.70
	ATOM	95	OE1	GLU	13	49.842	106.055	1.664	1.00	74.10
	ATOM	96	OE2	GLU	13	48.724	105.351	-0.125	1.00	71.55
	ATOM	97	C	GLU	13	46.341	106.579	5.195	1.00	43.24
35	ATOM	98	O	GLU	13	47.190	107.458	5.360	1.00	44.25
	ATOM	99	N	THR	14	45.207	106.548	5.879	1.00	37.06
	ATOM	100	CA	THR	14	44.959	107.523	6.934	1.00	36.77
	ATOM	101	CB	THR	14	44.105	108.735	6.457	1.00	33.39
	ATOM	102	OG1	THR	14	42.741	108.344	6.308	1.00	41.69
40	ATOM	103	CG2	THR	14	44.624	109.267	5.131	1.00	31.66
	ATOM	104	C	THR	14	44.243	106.773	8.052	1.00	34.96
	ATOM	105	O	THR	14	43.532	105.806	7.805	1.00	35.04
	ATOM	106	N	PRO	15	44.442	107.199	9.299	1.00	37.11
	ATOM	107	CD	PRO	15	45.296	108.325	9.705	1.00	36.50
45	ATOM	108	CA	PRO	15	43.823	106.564	10.468	1.00	38.81
	ATOM	109	CB	PRO	15	44.446	107.319	11.642	1.00	39.15
	ATOM	110	CG	PRO	15	45.719	107.895	11.071	1.00	37.32
	ATOM	111	C	PRO	15	42.305	106.687	10.477	1.00	37.78
	ATOM	112	O	PRO	15	41.759	107.640	9.929	1.00	35.91
50	ATOM	113	N	THR	16	41.620	105.733	11.099	1.00	36.67
	ATOM	114	CA	THR	16	40.170	105.815	11.164	1.00	38.46
	ATOM	115	CB	THR	16	39.542	104.541	11.760	1.00	38.61
	ATOM	116	OG1	THR	16	39.970	104.383	13.117	1.00	42.44
	ATOM	117	CG2	THR	16	39.957	103.320	10.955	1.00	40.11
55	ATOM	118	C	THR	16	39.865	106.979	12.082	1.00	35.97
	ATOM	119	O	THR	16	40.521	107.140	13.103	1.00	39.85
	ATOM	120	N	ILE	17	38.881	107.792	11.721	1.00	33.66
	ATOM	121	CA	ILE	17	38.522	108.946	12.532	1.00	34.76
	ATOM	122	CB	ILE	17	37.694	109.946	11.715	1.00	32.44
60	ATOM	123	CG2	ILE	17	37.224	111.082	12.602	1.00	28.66
	ATOM	124	CG1	ILE	17	38.533	110.464	10.541	1.00	33.15
	ATOM	125	CD1	ILE	17	37.795	111.404	9.609	1.00	31.19
	ATOM	126	C	ILE	17	37.747	108.584	13.792	1.00	40.64
	ATOM	127	O	ILE	17	36.743	107.886	13.734	1.00	44.07
65	ATOM	128	N	GLN	18	38.221	109.063	14.939	1.00	45.91
	ATOM	129	CA	GLN	18	37.559	108.798	16.214	1.00	50.22

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	ATOM	130	CB	GLN	18	38.575	108.396	17.278	1.00	53.09
	ATOM	131	CG	GLN	18	38.275	107.053	17.890	1.00	57.60
	ATOM	132	CD	GLN	18	38.479	105.945	16.890	1.00	61.39
	ATOM	133	OE1	GLN	18	39.610	105.653	16.493	1.00	68.36
5	ATOM	134	NE2	GLN	18	37.386	105.331	16.457	1.00	64.93
	ATOM	135	C	GLN	18	36.815	110.030	16.697	1.00	51.56
	ATOM	136	O	GLN	18	37.358	111.131	16.703	1.00	56.44
	ATOM	137	N	LYS	19	35.573	109.846	17.111	1.00	52.60
	ATOM	138	CA	LYS	19	34.792	110.973	17.592	1.00	56.10
10	ATOM	139	CB	LYS	19	34.447	111.918	16.440	1.00	55.55
	ATOM	140	CG	LYS	19	33.542	113.043	16.887	1.00	61.15
	ATOM	141	CD	LYS	19	33.359	114.115	15.833	1.00	64.42
	ATOM	142	CE	LYS	19	32.446	115.229	16.366	1.00	64.84
	ATOM	143	NZ	LYS	19	32.252	116.342	15.382	1.00	69.52
15	ATOM	144	C	LYS	19	33.511	110.543	18.291	1.00	57.41
	ATOM	145	O	LYS	19	32.786	109.679	17.794	1.00	61.02
	ATOM	146	N	GLY	20	33.239	111.159	19.441	1.00	58.84
	ATOM	147	CA	GLY	20	32.047	110.843	20.205	1.00	56.66
	ATOM	148	C	GLY	20	31.848	109.353	20.405	1.00	56.87
20	ATOM	149	O	GLY	20	30.713	108.875	20.368	1.00	58.64
	ATOM	150	N	SER	21	32.946	108.626	20.619	1.00	54.49
	ATOM	151	CA	SER	21	32.910	107.176	20.825	1.00	53.68
	ATOM	152	CB	SER	21	32.088	106.833	22.075	1.00	55.80
	ATOM	153	OG	SER	21	30.704	106.752	21.789	1.00	65.34
25	ATOM	154	C	SER	21	32.362	106.413	19.595	1.00	51.41
	ATOM	155	O	SER	21	31.804	105.308	19.709	1.00	47.74
	ATOM	156	N	TYR	22	32.528	107.025	18.423	1.00	46.25
	ATOM	157	CA	TYR	22	32.108	106.445	17.155	1.00	40.57
	ATOM	158	CB	TYR	22	31.061	107.331	16.476	1.00	43.27
30	ATOM	159	CG	TYR	22	29.642	107.041	16.892	1.00	49.18
	ATOM	160	CD1	TYR	22	29.248	107.148	18.221	1.00	52.61
	ATOM	161	CE1	TYR	22	27.936	106.867	18.613	1.00	51.92
	ATOM	162	CD2	TYR	22	28.690	106.647	15.958	1.00	50.01
	ATOM	163	CE2	TYR	22	27.377	106.367	16.338	1.00	50.94
35	ATOM	164	CZ	TYR	22	27.008	106.478	17.666	1.00	50.78
	ATOM	165	OH	TYR	22	25.713	106.198	18.042	1.00	53.87
	ATOM	166	C	TYR	22	33.339	106.351	16.260	1.00	37.85
	ATOM	167	O	TYR	22	34.251	107.170	16.368	1.00	38.70
	ATOM	168	N	THR	23	33.382	105.349	15.392	1.00	33.81
40	ATOM	169	CA	THR	23	34.510	105.209	14.478	1.00	32.40
	ATOM	170	CB	THR	23	35.071	103.765	14.436	1.00	31.58
	ATOM	171	OG1	THR	23	35.454	103.347	15.750	1.00	30.76
	ATOM	172	CG2	THR	23	36.279	103.703	13.526	1.00	24.08
	ATOM	173	C	THR	23	34.021	105.560	13.084	1.00	33.28
45	ATOM	174	O	THR	23	32.981	105.077	12.650	1.00	30.44
	ATOM	175	N	PHE	24	34.770	106.403	12.389	1.00	32.16
	ATOM	176	CA	PHE	24	34.404	106.800	11.041	1.00	30.68
	ATOM	177	CB	PHE	24	34.181	108.310	10.968	1.00	28.20
	ATOM	178	CG	PHE	24	33.002	108.782	11.758	1.00	31.00
50	ATOM	179	CD1	PHE	24	33.116	109.039	13.114	1.00	31.69
	ATOM	180	CD2	PHE	24	31.760	108.941	11.150	1.00	31.42
	ATOM	181	CE1	PHE	24	32.012	109.445	13.853	1.00	32.57
	ATOM	182	CE2	PHE	24	30.656	109.346	11.880	1.00	25.01
	ATOM	183	CZ	PHE	24	30.783	109.598	13.234	1.00	30.12
55	ATOM	184	C	PHE	24	35.462	106.388	10.025	1.00	31.73
	ATOM	185	O	PHE	24	36.648	106.646	10.203	1.00	33.31
	ATOM	186	N	VAL	25	35.021	105.737	8.961	1.00	32.92
	ATOM	187	CA	VAL	25	35.922	105.302	7.916	1.00	30.84
	ATOM	188	CB	VAL	25	35.201	104.385	6.915	1.00	30.29
60	ATOM	189	CG1	VAL	25	36.146	103.977	5.812	1.00	28.13
	ATOM	190	CG2	VAL	25	34.659	103.174	7.624	1.00	29.53
	ATOM	191	C	VAL	25	36.455	106.508	7.151	1.00	36.08
	ATOM	192	O	VAL	25	35.709	107.432	6.834	1.00	35.82
	ATOM	193	N	PRO	26	37.765	106.527	6.867	1.00	38.03
65	ATOM	194	CD	PRO	26	38.786	105.646	7.455	1.00	38.28
	ATOM	195	CA	PRO	26	38.400	107.626	6.126	1.00	38.08

	ATOM	196	CB	PRO	26	39.889	107.445	6.432	1.00	37.62
	ATOM	197	CG	PRO	26	39.916	106.601	7.666	1.00	36.31
	ATOM	198	C	PRO	26	38.108	107.362	4.650	1.00	40.85
	ATOM	199	O	PRO	26	38.621	106.400	4.080	1.00	42.27
5	ATOM	200	N	TRP	27	37.294	108.197	4.021	1.00	40.86
	ATOM	201	CA	TRP	27	36.956	107.949	2.625	1.00	37.05
	ATOM	202	CB	TRP	27	35.526	108.409	2.332	1.00	33.29
	ATOM	203	CG	TRP	27	34.500	107.711	3.150	1.00	31.06
	ATOM	204	CD2	TRP	27	34.222	106.310	3.163	1.00	27.79
10	ATOM	205	CE2	TRP	27	33.203	106.099	4.106	1.00	29.28
	ATOM	206	CE3	TRP	27	34.736	105.213	2.471	1.00	27.06
	ATOM	207	CD1	TRP	27	33.662	108.276	4.053	1.00	31.07
	ATOM	208	NE1	TRP	27	32.879	107.319	4.634	1.00	32.70
	ATOM	209	CZ2	TRP	27	32.683	104.831	4.380	1.00	25.99
15	ATOM	210	CZ3	TRP	27	34.219	103.951	2.743	1.00	26.93
	ATOM	211	CH2	TRP	27	33.203	103.773	3.690	1.00	28.08
	ATOM	212	C	TRP	27	37.887	108.562	1.604	1.00	38.50
	ATOM	213	O	TRP	27	38.602	109.519	1.881	1.00	39.07
	ATOM	214	N	LEU	28	37.858	107.975	0.413	1.00	40.19
20	ATOM	215	CA	LEU	28	38.648	108.420	-0.724	1.00	39.45
	ATOM	216	CB	LEU	28	39.822	107.477	-0.954	1.00	44.18
	ATOM	217	CG	LEU	28	40.940	108.013	-1.846	1.00	46.83
	ATOM	218	CD1	LEU	28	41.636	109.170	-1.123	1.00	45.15
	ATOM	219	CD2	LEU	28	41.934	106.892	-2.163	1.00	49.15
25	ATOM	220	C	LEU	28	37.684	108.323	-1.899	1.00	39.80
	ATOM	221	O	LEU	28	36.927	107.367	-1.990	1.00	42.40
	ATOM	222	N	LEU	29	37.692	109.300	-2.790	1.00	33.58
	ATOM	223	CA	LEU	29	36.781	109.244	-3.916	1.00	29.21
	ATOM	224	CB	LEU	29	36.840	110.535	-4.731	1.00	27.16
30	ATOM	225	CG	LEU	29	35.953	110.528	-5.980	1.00	22.36
	ATOM	226	CD1	LEU	29	34.496	110.666	-5.571	1.00	21.47
	ATOM	227	CD2	LEU	29	36.350	111.642	-6.911	1.00	14.53
	ATOM	228	C	LEU	29	37.069	108.069	-4.843	1.00	29.98
	ATOM	229	O	LEU	29	38.190	107.880	-5.305	1.00	30.36
35	ATOM	230	N	SER	30	36.037	107.278	-5.103	1.00	29.88
	ATOM	231	CA	SER	30	36.152	106.148	-6.005	1.00	28.49
	ATOM	232	CB	SER	30	35.159	105.048	-5.629	1.00	26.13
	ATOM	233	OG	SER	30	35.207	103.992	-6.562	1.00	20.25
	ATOM	234	C	SER	30	35.815	106.724	-7.374	1.00	31.61
40	ATOM	235	O	SER	30	36.576	106.581	-8.326	1.00	35.14
	ATOM	236	N	PHE	31	34.667	107.388	-7.460	1.00	32.12
	ATOM	237	CA	PHE	31	34.241	108.018	-8.698	1.00	30.76
	ATOM	238	CB	PHE	31	33.913	106.964	-9.763	1.00	27.55
	ATOM	239	CG	PHE	31	32.493	106.472	-9.731	1.00	31.23
45	ATOM	240	CD1	PHE	31	31.478	107.195	-10.348	1.00	32.00
	ATOM	241	CD2	PHE	31	32.165	105.290	-9.071	1.00	27.89
	ATOM	242	CE1	PHE	31	30.166	106.751	-10.309	1.00	29.70
	ATOM	243	CE2	PHE	31	30.855	104.842	-9.030	1.00	26.15
	ATOM	244	CZ	PHE	31	29.854	105.572	-9.648	1.00	28.30
50	ATOM	245	C	PHE	31	33.032	108.900	-8.439	1.00	32.06
	ATOM	246	O	PHE	31	32.242	108.651	-7.529	1.00	32.26
	ATOM	247	N	LYS	32	32.906	109.945	-9.238	1.00	34.83
	ATOM	248	CA	LYS	32	31.790	110.861	-9.118	1.00	37.13
	ATOM	249	CB	LYS	32	32.270	112.206	-8.576	1.00	36.55
55	ATOM	250	CG	LYS	32	31.179	113.236	-8.484	1.00	40.62
	ATOM	251	CD	LYS	32	31.725	114.617	-8.220	1.00	44.56
	ATOM	252	CE	LYS	32	30.616	115.648	-8.442	1.00	46.29
	ATOM	253	NZ	LYS	32	31.027	117.038	-8.084	1.00	50.58
	ATOM	254	C	LYS	32	31.201	111.025	-10.516	1.00	36.39
60	ATOM	255	O	LYS	32	31.935	111.213	-11.487	1.00	38.69
	ATOM	256	N	ARG	33	29.880	110.933	-10.622	1.00	37.90
	ATOM	257	CA	ARG	33	29.218	111.066	-11.912	1.00	35.74
	ATOM	258	CB	ARG	33	28.764	109.690	-12.386	1.00	36.79
	ATOM	259	CG	ARG	33	27.986	109.671	-13.682	1.00	31.09
65	ATOM	260	CD	ARG	33	27.841	108.235	-14.170	1.00	31.62
	ATOM	261	NE	ARG	33	27.233	108.131	-15.494	1.00	38.16

	ATOM	262	CZ	ARG	33	25.961	107.817	-15.709	1.00	39.15
	ATOM	263	NH1	ARG	33	25.155	107.571	-14.686	1.00	43.41
	ATOM	264	NH2	ARG	33	25.491	107.751	-16.943	1.00	40.65
	ATOM	265	C	ARG	33	28.027	112.010	-11.795	1.00	37.71
5	ATOM	266	O	ARG	33	27.092	111.756	-11.030	1.00	41.16
	ATOM	267	N	GLY	34	28.057	113.107	-12.543	1.00	36.59
	ATOM	268	CA	GLY	34	26.957	114.046	-12.480	1.00	36.72
	ATOM	269	C	GLY	34	27.186	115.148	-11.467	1.00	40.01
	ATOM	270	O	GLY	34	28.298	115.342	-10.977	1.00	39.85
10	ATOM	271	N	SER	35	26.115	115.855	-11.133	1.00	38.74
	ATOM	272	CA	SER	35	26.193	116.971	-10.202	1.00	40.23
	ATOM	273	CB	SER	35	25.744	118.236	-10.922	1.00	43.02
	ATOM	274	OG	SER	35	24.458	118.040	-11.509	1.00	48.05
	ATOM	275	C	SER	35	25.378	116.828	-8.914	1.00	40.54
15	ATOM	276	O	SER	35	25.626	117.552	-7.951	1.00	43.74
	ATOM	277	N	ALA	36	24.418	115.907	-8.894	1.00	35.56
	ATOM	278	CA	ALA	36	23.553	115.716	-7.735	1.00	29.62
	ATOM	279	CB	ALA	36	22.505	114.671	-8.056	1.00	25.47
	ATOM	280	C	ALA	36	24.244	115.365	-6.420	1.00	31.23
20	ATOM	281	O	ALA	36	23.704	115.634	-5.354	1.00	29.56
	ATOM	282	N	LEU	37	25.430	114.770	-6.485	1.00	34.09
	ATOM	283	CA	LEU	37	26.143	114.372	-5.275	1.00	32.32
	ATOM	284	CB	LEU	37	26.096	112.853	-5.125	1.00	26.47
	ATOM	285	CG	LEU	37	24.693	112.260	-5.026	1.00	25.72
25	ATOM	286	CD1	LEU	37	24.721	110.787	-5.349	1.00	24.61
	ATOM	287	CD2	LEU	37	24.134	112.505	-3.644	1.00	27.53
	ATOM	288	C	LEU	37	27.588	114.838	-5.254	1.00	35.16
	ATOM	289	O	LEU	37	28.269	114.840	-6.273	1.00	36.47
	ATOM	290	N	GLU	38	28.049	115.226	-4.075	1.00	38.30
30	ATOM	291	CA	GLU	38	29.414	115.694	-3.896	1.00	41.17
	ATOM	292	CB	GLU	38	29.444	117.214	-3.790	1.00	46.15
	ATOM	293	CG	GLU	38	29.569	117.957	-5.099	1.00	50.01
	ATOM	294	CD	GLU	38	29.310	119.448	-4.924	1.00	51.18
	ATOM	295	OE1	GLU	38	29.765	120.009	-3.894	1.00	48.40
35	ATOM	296	OE2	GLU	38	28.654	120.051	-5.814	1.00	50.23
	ATOM	297	C	GLU	38	30.022	115.131	-2.626	1.00	42.69
	ATOM	298	O	GLU	38	29.317	114.747	-1.702	1.00	42.55
	ATOM	299	N	GLU	39	31.343	115.084	-2.578	1.00	44.75
	ATOM	300	CA	GLU	39	32.011	114.598	-1.382	1.00	46.50
40	ATOM	301	CB	GLU	39	33.290	113.851	-1.747	1.00	47.52
	ATOM	302	CG	GLU	39	34.040	114.493	-2.894	1.00	58.50
	ATOM	303	CD	GLU	39	35.377	113.828	-3.159	1.00	62.35
	ATOM	304	OE1	GLU	39	36.004	114.139	-4.205	1.00	64.58
	ATOM	305	OE2	GLU	39	35.803	113.004	-2.316	1.00	63.04
45	ATOM	306	C	GLU	39	32.336	115.860	-0.605	1.00	44.56
	ATOM	307	O	GLU	39	32.753	116.855	-1.184	1.00	48.11
	ATOM	308	N	LYS	40	32.125	115.833	0.701	1.00	40.71
	ATOM	309	CA	LYS	40	32.395	116.998	1.520	1.00	38.08
	ATOM	310	CB	LYS	40	31.171	117.911	1.556	1.00	41.28
50	ATOM	311	CG	LYS	40	31.292	119.073	2.534	1.00	41.52
	ATOM	312	CD	LYS	40	29.940	119.734	2.764	1.00	42.73
	ATOM	313	CE	LYS	40	29.996	120.791	3.853	1.00	41.97
	ATOM	314	NZ	LYS	40	28.632	121.332	4.141	1.00	45.94
	ATOM	315	C	LYS	40	32.761	116.599	2.934	1.00	39.66
55	ATOM	316	O	LYS	40	31.940	116.062	3.673	1.00	36.18
	ATOM	317	N	GLU	41	34.008	116.866	3.300	1.00	40.16
	ATOM	318	CA	GLU	41	34.505	116.565	4.634	1.00	42.80
	ATOM	319	CB	GLU	41	33.952	117.597	5.610	1.00	45.31
	ATOM	320	CG	GLU	41	34.271	119.021	5.170	1.00	56.07
60	ATOM	321	CD	GLU	41	33.401	120.075	5.849	1.00	59.42
	ATOM	322	OE1	GLU	41	32.151	119.970	5.781	1.00	63.71
	ATOM	323	OE2	GLU	41	33.970	121.015	6.447	1.00	63.71
	ATOM	324	C	GLU	41	34.155	115.155	5.081	1.00	38.93
	ATOM	325	O	GLU	41	33.569	114.955	6.141	1.00	39.03
65	ATOM	326	N	ASN	42	34.507	114.187	4.246	1.00	34.69
	ATOM	327	CA	ASN	42	34.278	112.781	4.531	1.00	32.96

	ATOM	328	CB	ASN	42	34.907	112.411	5.869	1.00	32.65
	ATOM	329	CG	ASN	42	35.453	111.017	5.871	1.00	35.50
	ATOM	330	OD1	ASN	42	35.267	110.270	6.822	1.00	40.26
	ATOM	331	ND2	ASN	42	36.143	110.656	4.800	1.00	29.28
5	ATOM	332	C	ASN	42	32.821	112.345	4.537	1.00	31.44
	ATOM	333	O	ASN	42	32.488	111.289	5.066	1.00	22.78
	ATOM	334	N	LYS	43	31.959	113.154	3.942	1.00	33.26
	ATOM	335	CA	LYS	43	30.541	112.841	3.883	1.00	33.68
	ATOM	336	CB	LYS	43	29.764	113.723	4.860	1.00	33.55
10	ATOM	337	CG	LYS	43	30.061	113.435	6.310	1.00	40.49
	ATOM	338	CD	LYS	43	29.336	114.401	7.212	1.00	45.39
	ATOM	339	CE	LYS	43	29.854	115.812	7.015	1.00	52.57
	ATOM	340	NZ	LYS	43	29.223	116.756	7.973	1.00	62.06
	ATOM	341	C	LYS	43	30.028	113.075	2.481	1.00	32.05
15	ATOM	342	O	LYS	43	30.700	113.698	1.670	1.00	37.32
	ATOM	343	N	ILE	44	28.843	112.564	2.184	1.00	31.31
	ATOM	344	CA	ILE	44	28.280	112.774	0.866	1.00	29.52
	ATOM	345	CB	ILE	44	27.624	111.502	0.327	1.00	26.38
	ATOM	346	CG2	ILE	44	27.009	111.779	-1.029	1.00	26.09
20	ATOM	347	CG1	ILE	44	28.679	110.403	0.194	1.00	28.61
	ATOM	348	CD1	ILE	44	28.160	109.108	-0.351	1.00	21.51
	ATOM	349	C	ILE	44	27.257	113.893	0.948	1.00	31.60
	ATOM	350	O	ILE	44	26.303	113.824	1.720	1.00	34.50
	ATOM	351	N	LEU	45	27.478	114.937	0.161	1.00	31.44
25	ATOM	352	CA	LEU	45	26.587	116.087	0.133	1.00	31.40
	ATOM	353	CB	LEU	45	27.409	117.370	0.046	1.00	33.60
	ATOM	354	CG	LEU	45	26.616	118.665	-0.109	1.00	34.36
	ATOM	355	CD1	LEU	45	25.810	118.939	1.144	1.00	29.00
	ATOM	356	CD2	LEU	45	27.582	119.801	-0.386	1.00	32.42
30	ATOM	357	C	LEU	45	25.605	116.030	-1.034	1.00	27.80
	ATOM	358	O	LEU	45	25.993	115.829	-2.180	1.00	27.94
	ATOM	359	N	VAL	46	24.327	116.210	-0.725	1.00	28.00
	ATOM	360	CA	VAL	46	23.277	116.197	-1.733	1.00	31.53
	ATOM	361	CB	VAL	46	21.949	115.742	-1.105	1.00	27.89
35	ATOM	362	CG1	VAL	46	20.865	115.676	-2.153	1.00	26.10
	ATOM	363	CG2	VAL	46	22.135	114.400	-0.452	1.00	23.39
	ATOM	364	C	VAL	46	23.111	117.608	-2.319	1.00	33.94
	ATOM	365	O	VAL	46	22.888	118.569	-1.580	1.00	39.68
	ATOM	366	N	LYS	47	23.216	117.730	-3.640	1.00	33.59
40	ATOM	367	CA	LYS	47	23.084	119.030	-4.293	1.00	37.05
	ATOM	368	CB	LYS	47	24.252	119.259	-5.258	1.00	34.49
	ATOM	369	CG	LYS	47	25.613	118.972	-4.649	1.00	37.53
	ATOM	370	CD	LYS	47	25.858	119.807	-3.399	1.00	41.69
	ATOM	371	CE	LYS	47	26.622	121.077	-3.724	1.00	42.95
45	ATOM	372	NZ	LYS	47	26.051	121.769	-4.902	1.00	44.50
	ATOM	373	C	LYS	47	21.760	119.206	-5.038	1.00	37.34
	ATOM	374	O	LYS	47	21.418	120.310	-5.450	1.00	43.04
	ATOM	375	N	GLU	48	21.026	118.116	-5.210	1.00	36.99
	ATOM	376	CA	GLU	48	19.731	118.142	-5.884	1.00	37.31
50	ATOM	377	CB	GLU	48	19.833	117.566	-7.286	1.00	36.53
	ATOM	378	CG	GLU	48	20.609	118.394	-8.267	1.00	46.74
	ATOM	379	CD	GLU	48	20.694	117.713	-9.621	1.00	52.39
	ATOM	380	OE1	GLU	48	19.666	117.122	-10.053	1.00	51.33
	ATOM	381	OE2	GLU	48	21.781	117.772	-10.252	1.00	54.55
55	ATOM	382	C	GLU	48	18.803	117.257	-5.089	1.00	36.94
	ATOM	383	O	GLU	48	19.146	116.115	-4.813	1.00	40.21
	ATOM	384	N	THR	49	17.625	117.748	-4.727	1.00	36.34
	ATOM	385	CA	THR	49	16.739	116.900	-3.945	1.00	35.48
	ATOM	386	CB	THR	49	15.669	117.726	-3.178	1.00	31.16
60	ATOM	387	OG1	THR	49	14.449	117.736	-3.916	1.00	32.48
	ATOM	388	CG2	THR	49	16.134	119.147	-2.973	1.00	30.36
	ATOM	389	C	THR	49	16.067	115.855	-4.840	1.00	34.59
	ATOM	390	O	THR	49	15.858	116.078	-6.031	1.00	35.13
	ATOM	391	N	GLY	50	15.755	114.703	-4.252	1.00	32.49
65	ATOM	392	CA	GLY	50	15.118	113.638	-4.996	1.00	28.17
	ATOM	393	C	GLY	50	15.225	112.324	-4.256	1.00	29.99

	ATOM	394	O	GLY	50	15.477	112.306	-3.062	1.00	29.76
	ATOM	395	N	TYR	51	15.022	111.222	-4.969	1.00	31.68
	ATOM	396	CA	TYR	51	15.115	109.890	-4.382	1.00	32.88
	ATOM	397	CB	TYR	51	13.964	109.002	-4.868	1.00	35.97
5	ATOM	398	CG	TYR	51	12.635	109.407	-4.302	1.00	43.03
	ATOM	399	CD1	TYR	51	11.908	110.463	-4.859	1.00	44.15
	ATOM	400	CE1	TYR	51	10.717	110.904	-4.280	1.00	47.81
	ATOM	401	CD2	TYR	51	12.139	108.792	-3.155	1.00	47.08
	ATOM	402	CE2	TYR	51	10.953	109.223	-2.562	1.00	51.79
10	ATOM	403	CZ	TYR	51	10.246	110.283	-3.126	1.00	51.15
	ATOM	404	OH	TYR	51	9.098	110.741	-2.507	1.00	52.20
	ATOM	405	C	TYR	51	16.448	109.239	-4.726	1.00	31.97
	ATOM	406	O	TYR	51	16.827	109.148	-5.891	1.00	30.01
	ATOM	407	N	PHE	52	17.161	108.786	-3.701	1.00	31.50
15	ATOM	408	CA	PHE	52	18.455	108.160	-3.913	1.00	30.41
	ATOM	409	CB	PHE	52	19.564	108.987	-3.259	1.00	31.30
	ATOM	410	CG	PHE	52	19.668	110.391	-3.767	1.00	29.70
	ATOM	411	CD1	PHE	52	18.728	111.344	-3.413	1.00	26.20
	ATOM	412	CD2	PHE	52	20.718	110.762	-4.594	1.00	28.38
20	ATOM	413	CE1	PHE	52	18.833	112.640	-3.875	1.00	27.17
	ATOM	414	CE2	PHE	52	20.830	112.058	-5.061	1.00	27.76
	ATOM	415	CZ	PHE	52	19.888	112.998	-4.702	1.00	27.27
	ATOM	416	C	PHE	52	18.545	106.751	-3.365	1.00	27.65
	ATOM	417	O	PHE	52	17.928	106.421	-2.360	1.00	24.85
25	ATOM	418	N	PHE	53	19.322	105.926	-4.050	1.00	27.18
	ATOM	419	CA	PHE	53	19.580	104.561	-3.620	1.00	27.05
	ATOM	420	CB	PHE	53	19.740	103.633	-4.817	1.00	28.76
	ATOM	421	CG	PHE	53	20.196	102.257	-4.453	1.00	28.00
	ATOM	422	CD1	PHE	53	19.350	101.379	-3.793	1.00	26.49
30	ATOM	423	CD2	PHE	53	21.485	101.840	-4.754	1.00	28.04
	ATOM	424	CE1	PHE	53	19.781	100.105	-3.440	1.00	26.54
	ATOM	425	CE2	PHE	53	21.920	100.566	-4.401	1.00	25.78
	ATOM	426	CZ	PHE	53	21.065	99.701	-3.744	1.00	24.39
	ATOM	427	C	PHE	53	20.918	104.700	-2.899	1.00	27.56
35	ATOM	428	O	PHE	53	21.898	105.136	-3.495	1.00	25.39
	ATOM	429	N	ILE	54	20.950	104.347	-1.618	1.00	28.87
	ATOM	430	CA	ILE	54	22.163	104.474	-0.817	1.00	28.95
	ATOM	431	CB	ILE	54	21.888	105.374	0.405	1.00	29.60
	ATOM	432	CG2	ILE	54	23.197	105.740	1.086	1.00	27.42
40	ATOM	433	CG1	ILE	54	21.163	106.643	-0.050	1.00	26.41
	ATOM	434	CD1	ILE	54	20.549	107.442	1.058	1.00	28.70
	ATOM	435	C	ILE	54	22.672	103.110	-0.352	1.00	28.57
	ATOM	436	O	ILE	54	21.898	102.259	0.073	1.00	29.17
	ATOM	437	N	TYR	55	23.980	102.900	-0.426	1.00	26.97
45	ATOM	438	CA	TYR	55	24.542	101.614	-0.031	1.00	24.22
	ATOM	439	CB	TYR	55	24.718	100.732	-1.268	1.00	21.82
	ATOM	440	CG	TYR	55	25.609	101.325	-2.331	1.00	27.68
	ATOM	441	CD1	TYR	55	26.979	101.081	-2.337	1.00	27.01
	ATOM	442	CE1	TYR	55	27.798	101.643	-3.300	1.00	27.44
50	ATOM	443	CD2	TYR	55	25.085	102.148	-3.322	1.00	27.16
	ATOM	444	CE2	TYR	55	25.900	102.717	-4.288	1.00	25.39
	ATOM	445	CZ	TYR	55	27.251	102.462	-4.271	1.00	29.02
	ATOM	446	OH	TYR	55	28.057	103.035	-5.221	1.00	31.82
	ATOM	447	C	TYR	55	25.856	101.739	0.709	1.00	25.16
55	ATOM	448	O	TYR	55	26.494	102.767	0.671	1.00	28.47
	ATOM	449	N	GLY	56	26.254	100.683	1.398	1.00	27.44
	ATOM	450	CA	GLY	56	27.499	100.723	2.134	1.00	22.07
	ATOM	451	C	GLY	56	27.911	99.357	2.637	1.00	25.83
	ATOM	452	O	GLY	56	27.066	98.531	2.967	1.00	26.00
60	ATOM	453	N	GLN	57	29.215	99.114	2.671	1.00	22.56
	ATOM	454	CA	GLN	57	29.758	97.856	3.157	1.00	20.91
	ATOM	455	CB	GLN	57	30.066	96.890	2.012	1.00	17.87
	ATOM	456	CG	GLN	57	30.759	95.619	2.488	1.00	16.86
	ATOM	457	CD	GLN	57	31.001	94.607	1.388	1.00	22.06
65	ATOM	458	OE1	GLN	57	30.063	94.072	0.807	1.00	27.26
	ATOM	459	NE2	GLN	57	32.268	94.335	1.101	1.00	20.20

	ATOM	460	C	GLN	57	31.033	98.134	3.925	1.00	24.23
	ATOM	461	O	GLN	57	31.791	99.035	3.570	1.00	25.16
	ATOM	462	N	VAL	58	31.255	97.353	4.978	1.00	24.62
	ATOM	463	CA	VAL	58	32.435	97.475	5.826	1.00	26.68
5	ATOM	464	CB	VAL	58	32.112	98.229	7.139	1.00	25.55
	ATOM	465	CG1	VAL	58	33.287	98.158	8.083	1.00	15.69
	ATOM	466	CG2	VAL	58	31.768	99.674	6.848	1.00	25.77
	ATOM	467	C	VAL	58	32.883	96.076	6.195	1.00	31.06
	ATOM	468	O	VAL	58	32.045	95.221	6.448	1.00	27.74
10	ATOM	469	N	LEU	59	34.195	95.838	6.220	1.00	33.82
	ATOM	470	CA	LEU	59	34.735	94.531	6.611	1.00	29.85
	ATOM	471	CB	LEU	59	35.933	94.142	5.750	1.00	29.62
	ATOM	472	CG	LEU	59	36.267	92.648	5.640	1.00	28.11
	ATOM	473	CD1	LEU	59	37.657	92.491	5.058	1.00	26.67
15	ATOM	474	CD2	LEU	59	36.196	91.968	6.975	1.00	22.45
	ATOM	475	C	LEU	59	35.188	94.630	8.065	1.00	30.98
	ATOM	476	O	LEU	59	36.119	95.366	8.385	1.00	32.52
	ATOM	477	N	TYR	60	34.523	93.889	8.945	1.00	33.01
	ATOM	478	CA	TYR	60	34.869	93.908	10.359	1.00	34.51
20	ATOM	479	CB	TYR	60	33.611	93.754	11.208	1.00	40.10
	ATOM	480	CG	TYR	60	32.641	94.876	11.004	1.00	45.37
	ATOM	481	CD1	TYR	60	31.469	94.690	10.270	1.00	47.13
	ATOM	482	CE1	TYR	60	30.616	95.754	10.011	1.00	50.28
	ATOM	483	CD2	TYR	60	32.930	96.149	11.478	1.00	44.35
25	ATOM	484	CE2	TYR	60	32.092	97.217	11.228	1.00	48.81
	ATOM	485	CZ	TYR	60	30.942	97.017	10.492	1.00	51.91
	ATOM	486	OH	TYR	60	30.139	98.095	10.213	1.00	53.77
	ATOM	487	C	TYR	60	35.875	92.837	10.745	1.00	35.49
	ATOM	488	O	TYR	60	35.652	91.649	10.538	1.00	36.13
30	ATOM	489	N	THR	61	36.991	93.277	11.308	1.00	36.19
	ATOM	490	CA	THR	61	38.037	92.374	11.748	1.00	35.36
	ATOM	491	CB	THR	61	39.342	92.646	10.998	1.00	34.22
	ATOM	492	OG1	THR	61	39.674	94.033	11.108	1.00	34.22
	ATOM	493	CG2	THR	61	39.200	92.274	9.534	1.00	25.45
35	ATOM	494	C	THR	61	38.230	92.619	13.238	1.00	37.82
	ATOM	495	O	THR	61	39.329	92.507	13.772	1.00	43.70
	ATOM	496	N	ASP	62	37.129	92.962	13.893	1.00	37.64
	ATOM	497	CA	ASP	62	37.087	93.251	15.315	1.00	38.08
	ATOM	498	CB	ASP	62	36.332	94.563	15.504	1.00	38.42
40	ATOM	499	CG	ASP	62	36.376	95.069	16.917	1.00	41.25
	ATOM	500	OD1	ASP	62	36.501	96.304	17.093	1.00	38.28
	ATOM	501	OD2	ASP	62	36.274	94.235	17.840	1.00	41.37
	ATOM	502	C	ASP	62	36.352	92.096	15.989	1.00	40.90
	ATOM	503	O	ASP	62	35.388	91.586	15.435	1.00	46.73
45	ATOM	504	N	LYS	63	36.788	91.671	17.172	1.00	42.62
	ATOM	505	CA	LYS	63	36.115	90.550	17.829	1.00	40.92
	ATOM	506	CB	LYS	63	37.144	89.587	18.424	1.00	42.79
	ATOM	507	CG	LYS	63	38.078	90.201	19.444	1.00	45.91
	ATOM	508	CD	LYS	63	39.191	89.217	19.805	1.00	46.22
50	ATOM	509	CE	LYS	63	40.046	88.871	18.571	1.00	49.97
	ATOM	510	NZ	LYS	63	41.016	87.744	18.796	1.00	49.98
	ATOM	511	C	LYS	63	35.095	90.931	18.897	1.00	42.59
	ATOM	512	O	LYS	63	34.731	90.109	19.728	1.00	42.46
	ATOM	513	N	THR	64	34.608	92.164	18.840	1.00	44.06
55	ATOM	514	CA	THR	64	33.636	92.665	19.801	1.00	44.94
	ATOM	515	CB	THR	64	33.490	94.201	19.662	1.00	43.79
	ATOM	516	OG1	THR	64	34.778	94.811	19.791	1.00	46.20
	ATOM	517	CG2	THR	64	32.579	94.768	20.745	1.00	42.80
	ATOM	518	C	THR	64	32.238	92.026	19.755	1.00	45.82
60	ATOM	519	O	THR	64	31.242	92.723	19.828	1.00	53.10
	ATOM	520	N	TYR	65	32.161	90.709	19.647	1.00	45.11
	ATOM	521	CA	TYR	65	30.885	89.978	19.633	1.00	45.72
	ATOM	522	CB	TYR	65	30.304	89.905	21.058	1.00	40.46
	ATOM	523	CG	TYR	65	29.306	90.982	21.421	1.00	41.90
65	ATOM	524	CD1	TYR	65	27.946	90.828	21.145	1.00	42.47
	ATOM	525	CE1	TYR	65	27.016	91.831	21.472	1.00	41.83

	ATOM	526	CD2	TYR	65	29.720	92.168	22.037	1.00	41.65
	ATOM	527	CE2	TYR	65	28.802	93.180	22.367	1.00	40.28
	ATOM	528	CZ	TYR	65	27.455	93.001	22.080	1.00	41.90
	ATOM	529	OH	TYR	65	26.554	93.986	22.400	1.00	43.84
5	ATOM	530	C	TYR	65	29.795	90.450	18.657	1.00	45.31
	ATOM	531	O	TYR	65	28.989	89.641	18.178	1.00	49.74
	ATOM	532	N	ALA	66	29.757	91.743	18.367	1.00	42.32
	ATOM	533	CA	ALA	66	28.764	92.293	17.454	1.00	39.57
	ATOM	534	CB	ALA	66	27.449	92.500	18.173	1.00	35.36
10	ATOM	535	C	ALA	66	29.262	93.610	16.887	1.00	39.85
	ATOM	536	O	ALA	66	29.560	94.547	17.630	1.00	37.60
	ATOM	537	N	MET	67	29.369	93.671	15.566	1.00	41.88
	ATOM	538	CA	MET	67	29.828	94.878	14.895	1.00	41.48
	ATOM	539	CB	MET	67	31.172	94.639	14.196	1.00	39.98
15	ATOM	540	CG	MET	67	32.352	94.420	15.134	1.00	37.31
	ATOM	541	SD	MET	67	32.637	95.805	16.244	1.00	35.46
	ATOM	542	CE	MET	67	33.538	96.941	15.187	1.00	34.06
	ATOM	543	C	MET	67	28.787	95.310	13.877	1.00	39.50
	ATOM	544	O	MET	67	27.877	94.549	13.541	1.00	38.44
20	ATOM	545	N	GLY	68	28.925	96.537	13.391	1.00	39.84
	ATOM	546	CA	GLY	68	27.987	97.048	12.412	1.00	37.61
	ATOM	547	C	GLY	68	28.172	98.526	12.166	1.00	33.76
	ATOM	548	O	GLY	68	28.947	99.180	12.855	1.00	35.58
	ATOM	549	N	HIS	69	27.471	99.054	11.172	1.00	35.07
25	ATOM	550	CA	HIS	69	27.564	100.469	10.855	1.00	33.70
	ATOM	551	CB	HIS	69	28.479	100.702	9.638	1.00	37.02
	ATOM	552	CG	HIS	69	28.114	99.903	8.425	1.00	34.65
	ATOM	553	CD2	HIS	69	27.522	100.268	7.265	1.00	35.63
	ATOM	554	ND1	HIS	69	28.375	98.555	8.311	1.00	37.68
30	ATOM	555	CE1	HIS	69	27.961	98.123	7.134	1.00	32.64
	ATOM	556	NE2	HIS	69	27.440	99.144	6.479	1.00	35.90
	ATOM	557	C	HIS	69	26.192	101.077	10.601	1.00	33.98
	ATOM	558	O	HIS	69	25.204	100.369	10.488	1.00	30.94
	ATOM	559	N	LEU	70	26.141	102.399	10.519	1.00	35.35
35	ATOM	560	CA	LEU	70	24.893	103.108	10.279	1.00	30.76
	ATOM	561	CB	LEU	70	24.523	103.954	11.492	1.00	29.59
	ATOM	562	CG	LEU	70	24.747	103.386	12.890	1.00	30.71
	ATOM	563	CD1	LEU	70	24.556	104.485	13.907	1.00	25.60
	ATOM	564	CD2	LEU	70	23.798	102.238	13.140	1.00	27.39
40	ATOM	565	C	LEU	70	25.046	104.056	9.094	1.00	33.19
	ATOM	566	O	LEU	70	26.072	104.721	8.952	1.00	37.07
	ATOM	567	N	ILE	71	24.042	104.104	8.226	1.00	30.93
	ATOM	568	CA	ILE	71	24.079	105.047	7.120	1.00	27.94
	ATOM	569	CB	ILE	71	23.529	104.448	5.823	1.00	28.22
45	ATOM	570	CG2	ILE	71	23.268	105.544	4.811	1.00	24.54
	ATOM	571	CG1	ILE	71	24.538	103.459	5.252	1.00	26.34
	ATOM	572	CD1	ILE	71	24.014	102.668	4.086	1.00	37.50
	ATOM	573	C	ILE	71	23.142	106.124	7.642	1.00	30.32
	ATOM	574	O	ILE	71	21.960	105.868	7.841	1.00	23.07
50	ATOM	575	N	GLN	72	23.673	107.317	7.895	1.00	29.74
	ATOM	576	CA	GLN	72	22.848	108.382	8.446	1.00	27.85
	ATOM	577	CB	GLN	72	23.449	108.848	9.763	1.00	26.70
	ATOM	578	CG	GLN	72	23.750	107.708	10.693	1.00	28.28
	ATOM	579	CD	GLN	72	24.155	108.177	12.061	1.00	33.11
55	ATOM	580	OE1	GLN	72	25.073	108.980	12.201	1.00	41.22
	ATOM	581	NE2	GLN	72	23.477	107.677	13.085	1.00	30.18
	ATOM	582	C	GLN	72	22.624	109.577	7.544	1.00	29.85
	ATOM	583	O	GLN	72	23.401	109.839	6.624	1.00	28.20
	ATOM	584	N	ARG	73	21.554	110.307	7.837	1.00	30.90
60	ATOM	585	CA	ARG	73	21.190	111.493	7.080	1.00	30.79
	ATOM	586	CB	ARG	73	19.854	111.261	6.388	1.00	29.18
	ATOM	587	CG	ARG	73	19.336	112.456	5.647	1.00	28.66
	ATOM	588	CD	ARG	73	17.848	112.360	5.488	1.00	29.25
	ATOM	589	NE	ARG	73	17.316	113.480	4.727	1.00	35.78
65	ATOM	590	CZ	ARG	73	16.042	113.841	4.728	1.00	35.27
	ATOM	591	NH1	ARG	73	15.162	113.172	5.459	1.00	34.95

	ATOM	592	NH2	ARG	73	15.651	114.867	3.994	1.00	33.33
	ATOM	593	C	ARG	73	21.081	112.727	7.973	1.00	32.42
	ATOM	594	O	ARG	73	20.397	112.698	8.990	1.00	34.19
	ATOM	595	N	LYS	74	21.773	113.798	7.597	1.00	35.90
5	ATOM	596	CA	LYS	74	21.708	115.057	8.339	1.00	38.57
	ATOM	597	CB	LYS	74	23.087	115.714	8.447	1.00	43.52
	ATOM	598	CG	LYS	74	24.045	115.032	9.409	1.00	52.89
	ATOM	599	CD	LYS	74	25.432	115.716	9.436	1.00	58.14
	ATOM	600	CE	LYS	74	26.377	114.998	10.401	1.00	56.75
10	ATOM	601	NZ	LYS	74	27.740	115.608	10.417	1.00	63.98
	ATOM	602	C	LYS	74	20.789	115.989	7.559	1.00	40.11
	ATOM	603	O	LYS	74	21.175	116.510	6.505	1.00	37.55
	ATOM	604	N	LYS	75	19.581	116.197	8.072	1.00	34.48
	ATOM	605	CA	LYS	75	18.607	117.059	7.413	1.00	34.24
15	ATOM	606	CB	LYS	75	17.251	116.945	8.114	1.00	36.16
	ATOM	607	CG	LYS	75	16.646	115.546	8.208	1.00	35.99
	ATOM	608	CD	LYS	75	15.305	115.617	8.950	1.00	40.91
	ATOM	609	CE	LYS	75	14.634	114.254	9.055	1.00	46.16
	ATOM	610	NZ	LYS	75	13.340	114.251	9.823	1.00	46.03
20	ATOM	611	C	LYS	75	19.048	118.526	7.433	1.00	30.92
	ATOM	612	O	LYS	75	19.570	118.995	8.427	1.00	27.47
	ATOM	613	N	VAL	76	18.828	119.242	6.333	1.00	32.82
	ATOM	614	CA	VAL	76	19.174	120.664	6.250	1.00	31.51
	ATOM	615	CB	VAL	76	19.192	121.197	4.806	1.00	29.19
25	ATOM	616	CG1	VAL	76	20.265	122.238	4.663	1.00	27.09
	ATOM	617	CG2	VAL	76	19.378	120.086	3.835	1.00	34.94
	ATOM	618	C	VAL	76	18.076	121.441	6.947	1.00	33.40
	ATOM	619	O	VAL	76	18.334	122.410	7.653	1.00	32.92
	ATOM	620	N	HIS	77	16.841	121.008	6.714	1.00	34.88
30	ATOM	621	CA	HIS	77	15.667	121.635	7.293	1.00	34.59
	ATOM	622	CB	HIS	77	14.607	121.865	6.211	1.00	35.92
	ATOM	623	CG	HIS	77	15.082	122.707	5.067	1.00	34.17
	ATOM	624	CD2	HIS	77	14.672	122.773	3.780	1.00	31.44
	ATOM	625	ND1	HIS	77	16.086	123.645	5.196	1.00	35.05
35	ATOM	626	CE1	HIS	77	16.277	124.251	4.040	1.00	29.84
	ATOM	627	NE2	HIS	77	15.430	123.742	3.165	1.00	38.35
	ATOM	628	C	HIS	77	15.119	120.745	8.396	1.00	35.63
	ATOM	629	O	HIS	77	15.073	119.524	8.245	1.00	40.24
	ATOM	630	N	VAL	78	14.680	121.356	9.495	1.00	37.37
40	ATOM	631	CA	VAL	78	14.193	120.577	10.619	1.00	41.32
	ATOM	632	CB	VAL	78	15.103	120.805	11.842	1.00	43.10
	ATOM	633	CG1	VAL	78	14.692	119.892	12.971	1.00	45.01
	ATOM	634	CG2	VAL	78	16.545	120.509	11.471	1.00	44.98
	ATOM	635	C	VAL	78	12.723	120.669	11.048	1.00	43.02
45	ATOM	636	O	VAL	78	11.999	119.683	10.920	1.00	49.79
	ATOM	637	N	PHE	79	12.259	121.800	11.563	1.00	41.33
	ATOM	638	CA	PHE	79	10.844	121.890	11.999	1.00	45.00
	ATOM	639	CB	PHE	79	9.868	121.214	11.015	1.00	40.51
	ATOM	640	CG	PHE	79	9.964	121.707	9.611	1.00	39.52
50	ATOM	641	CD1	PHE	79	10.575	120.929	8.634	1.00	36.80
	ATOM	642	CD2	PHE	79	9.440	122.944	9.260	1.00	37.89
	ATOM	643	CE1	PHE	79	10.663	121.375	7.326	1.00	37.52
	ATOM	644	CE2	PHE	79	9.523	123.397	7.953	1.00	35.59
	ATOM	645	CZ	PHE	79	10.136	122.612	6.983	1.00	37.58
55	ATOM	646	C	PHE	79	10.555	121.248	13.356	1.00	43.50
	ATOM	647	O	PHE	79	10.818	120.063	13.569	1.00	37.85
	ATOM	648	N	GLY	80	9.974	122.037	14.253	1.00	46.65
	ATOM	649	CA	GLY	80	9.605	121.554	15.569	1.00	48.38
	ATOM	650	C	GLY	80	10.628	120.738	16.328	1.00	47.89
60	ATOM	651	O	GLY	80	11.795	121.125	16.437	1.00	51.15
	ATOM	652	N	ASP	81	10.184	119.599	16.855	1.00	45.07
	ATOM	653	CA	ASP	81	11.048	118.734	17.637	1.00	43.99
	ATOM	654	CB	ASP	81	10.281	118.178	18.844	1.00	45.66
	ATOM	655	CG	ASP	81	9.198	117.185	18.452	1.00	47.49
65	ATOM	656	OD1	ASP	81	8.842	117.105	17.255	1.00	51.90
	ATOM	657	OD2	ASP	81	8.690	116.484	19.349	1.00	48.08

	ATOM	658	C	ASP	81	11.671	117.596	16.848	1.00	42.29
	ATOM	659	O	ASP	81	12.053	116.578	17.424	1.00	39.94
	ATOM	660	N	GLU	82	11.773	117.763	15.533	1.00	39.91
	ATOM	661	CA	GLU	82	12.391	116.740	14.692	1.00	36.58
5	ATOM	662	CB	GLU	82	12.346	117.120	13.211	1.00	34.91
	ATOM	663	CG	GLU	82	11.058	116.907	12.483	1.00	37.82
	ATOM	664	CD	GLU	82	11.289	116.739	10.990	1.00	40.75
	ATOM	665	OE1	GLU	82	12.093	117.501	10.417	1.00	38.15
	ATOM	666	OE2	GLU	82	10.671	115.840	10.382	1.00	47.99
10	ATOM	667	C	GLU	82	13.864	116.616	15.054	1.00	36.58
	ATOM	668	O	GLU	82	14.495	117.590	15.456	1.00	38.40
	ATOM	669	N	LEU	83	14.408	115.416	14.911	1.00	33.83
	ATOM	670	CA	LEU	83	15.825	115.192	15.147	1.00	32.70
	ATOM	671	CB	LEU	83	16.086	113.743	15.557	1.00	30.78
15	ATOM	672	CG	LEU	83	16.119	113.415	17.048	1.00	29.26
	ATOM	673	CD1	LEU	83	14.999	114.113	17.778	1.00	29.60
	ATOM	674	CD2	LEU	83	16.021	111.921	17.222	1.00	28.65
	ATOM	675	C	LEU	83	16.405	115.450	13.773	1.00	33.88
	ATOM	676	O	LEU	83	15.879	114.950	12.785	1.00	38.46
20	ATOM	677	N	SER	84	17.470	116.235	13.693	1.00	34.94
	ATOM	678	CA	SER	84	18.054	116.536	12.395	1.00	39.77
	ATOM	679	CB	SER	84	18.834	117.848	12.453	1.00	38.92
	ATOM	680	OG	SER	84	19.852	117.774	13.421	1.00	43.11
	ATOM	681	C	SER	84	18.949	115.421	11.862	1.00	39.89
25	ATOM	682	O	SER	84	19.348	115.430	10.698	1.00	42.52
	ATOM	683	N	LEU	85	19.257	114.457	12.717	1.00	40.92
	ATOM	684	CA	LEU	85	20.088	113.335	12.325	1.00	34.66
	ATOM	685	CB	LEU	85	21.243	113.171	13.313	1.00	35.12
	ATOM	686	CG	LEU	85	22.432	112.247	13.013	1.00	36.92
30	ATOM	687	CD1	LEU	85	21.990	110.804	13.022	1.00	38.24
	ATOM	688	CD2	LEU	85	23.048	112.620	11.673	1.00	36.08
	ATOM	689	C	LEU	85	19.204	112.102	12.327	1.00	35.30
	ATOM	690	O	LEU	85	18.689	111.690	13.363	1.00	32.85
	ATOM	691	N	VAL	86	19.008	111.528	11.150	1.00	33.26
35	ATOM	692	CA	VAL	86	18.185	110.337	11.013	1.00	34.57
	ATOM	693	CB	VAL	86	17.056	110.531	9.960	1.00	35.39
	ATOM	694	CG1	VAL	86	16.331	109.221	9.719	1.00	31.44
	ATOM	695	CG2	VAL	86	16.075	111.584	10.436	1.00	38.08
	ATOM	696	C	VAL	86	19.052	109.190	10.551	1.00	32.85
40	ATOM	697	O	VAL	86	19.850	109.342	9.630	1.00	35.10
	ATOM	698	N	THR	87	18.917	108.041	11.189	1.00	33.41
	ATOM	699	CA	THR	87	19.690	106.907	10.743	1.00	33.56
	ATOM	700	CB	THR	87	20.378	106.172	11.935	1.00	32.70
	ATOM	701	OG1	THR	87	20.307	104.753	11.748	1.00	34.29
45	ATOM	702	CG2	THR	87	19.755	106.576	13.245	1.00	34.95
	ATOM	703	C	THR	87	18.810	105.965	9.912	1.00	30.57
	ATOM	704	O	THR	87	17.830	105.407	10.399	1.00	28.08
	ATOM	705	N	LEU	88	19.148	105.864	8.630	1.00	28.42
	ATOM	706	CA	LEU	88	18.462	104.990	7.697	1.00	31.31
50	ATOM	707	CB	LEU	88	18.516	105.553	6.270	1.00	28.21
	ATOM	708	CG	LEU	88	18.117	106.938	5.751	1.00	29.12
	ATOM	709	CD1	LEU	88	17.430	107.738	6.801	1.00	31.12
	ATOM	710	CD2	LEU	88	19.351	107.651	5.253	1.00	28.72
	ATOM	711	C	LEU	88	19.263	103.677	7.693	1.00	41.15
55	ATOM	712	O	LEU	88	20.482	103.674	7.884	1.00	52.30
	ATOM	713	N	PHE	89	18.603	102.550	7.502	1.00	39.23
	ATOM	714	CA	PHE	89	19.350	101.294	7.403	1.00	42.27
	ATOM	715	CB	PHE	89	20.233	101.384	6.155	1.00	35.63
	ATOM	716	CG	PHE	89	19.520	102.021	4.991	1.00	35.06
60	ATOM	717	CD1	PHE	89	20.157	102.970	4.198	1.00	31.56
	ATOM	718	CD2	PHE	89	18.157	101.757	4.768	1.00	30.96
	ATOM	719	CE1	PHE	89	19.446	103.661	3.211	1.00	33.76
	ATOM	720	CE2	PHE	89	17.441	102.439	3.791	1.00	27.38
	ATOM	721	CZ	PHE	89	18.083	103.395	3.013	1.00	30.29
65	ATOM	722	C	PHE	89	20.139	100.785	8.630	1.00	39.99
	ATOM	723	O	PHE	89	19.536	100.481	9.664	1.00	45.34

	ATOM	724	N	ARG	90	21.455	100.657	8.546	1.00	38.92
	ATOM	725	CA	ARG	90	22.200	100.110	9.695	1.00	41.66
	ATOM	726	CB	ARG	90	21.673	100.692	11.023	1.00	39.24
	ATOM	727	CG	ARG	90	21.654	99.702	12.205	1.00	29.41
5	ATOM	728	CD	ARG	90	21.145	100.351	13.506	1.00	31.13
	ATOM	729	NE	ARG	90	20.865	99.361	14.549	1.00	31.75
	ATOM	730	CZ	ARG	90	19.653	99.083	15.036	1.00	28.86
	ATOM	731	NH1	ARG	90	18.579	99.718	14.597	1.00	25.48
	ATOM	732	NH2	ARG	90	19.509	98.138	15.950	1.00	29.84
10	ATOM	733	C	ARG	90	22.158	98.574	9.808	1.00	39.23
	ATOM	734	O	ARG	90	21.088	97.969	9.768	1.00	31.33
	ATOM	735	N	CYS	91	23.330	97.950	9.948	1.00	40.12
	ATOM	736	CA	CYS	91	23.391	96.506	10.113	1.00	37.09
	ATOM	737	C	CYS	91	24.324	95.990	11.171	1.00	37.21
15	ATOM	738	O	CYS	91	25.162	96.719	11.687	1.00	37.46
	ATOM	739	CB	CYS	91	23.703	95.793	8.818	1.00	40.13
	ATOM	740	SG	CYS	91	25.259	96.056	7.895	1.00	41.64
	ATOM	741	N	ILE	92	24.164	94.708	11.481	1.00	34.02
	ATOM	742	CA	ILE	92	24.954	94.056	12.515	1.00	33.22
20	ATOM	743	CB	ILE	92	24.110	93.880	13.794	1.00	32.88
	ATOM	744	CG2	ILE	92	24.979	93.382	14.929	1.00	32.61
	ATOM	745	CG1	ILE	92	23.472	95.217	14.178	1.00	34.64
	ATOM	746	CD1	ILE	92	22.461	95.129	15.310	1.00	32.30
	ATOM	747	C	ILE	92	25.470	92.686	12.093	1.00	30.78
25	ATOM	748	O	ILE	92	24.880	92.030	11.260	1.00	33.43
	ATOM	749	N	GLN	93	26.597	92.275	12.662	1.00	29.35
	ATOM	750	CA	GLN	93	27.182	90.973	12.386	1.00	28.69
	ATOM	751	CB	GLN	93	28.296	91.056	11.342	1.00	28.17
	ATOM	752	CG	GLN	93	27.880	91.028	9.869	1.00	25.87
30	ATOM	753	CD	GLN	93	27.151	89.766	9.462	1.00	24.22
	ATOM	754	OE1	GLN	93	25.937	89.686	9.566	1.00	28.90
	ATOM	755	NE2	GLN	93	27.894	88.771	8.997	1.00	25.29
	ATOM	756	C	GLN	93	27.792	90.475	13.677	1.00	31.44
	ATOM	757	O	GLN	93	28.576	91.179	14.295	1.00	32.07
35	ATOM	758	N	ASN	94	27.416	89.280	14.110	1.00	35.77
	ATOM	759	CA	ASN	94	28.011	88.726	15.311	1.00	33.56
	ATOM	760	CB	ASN	94	27.266	87.469	15.764	1.00	34.70
	ATOM	761	CG	ASN	94	26.020	87.782	16.553	1.00	33.01
	ATOM	762	OD1	ASN	94	26.061	88.537	17.510	1.00	30.56
40	ATOM	763	ND2	ASN	94	24.909	87.188	16.166	1.00	36.61
	ATOM	764	C	ASN	94	29.446	88.376	14.917	1.00	35.80
	ATOM	765	O	ASN	94	29.702	87.944	13.791	1.00	36.63
	ATOM	766	N	MET	95	30.385	88.572	15.833	1.00	35.73
	ATOM	767	CA	MET	95	31.784	88.281	15.544	1.00	34.73
45	ATOM	768	CB	MET	95	32.641	89.509	15.865	1.00	32.97
	ATOM	769	CG	MET	95	32.201	90.797	15.184	1.00	27.35
	ATOM	770	SD	MET	95	32.070	90.670	13.389	1.00	25.14
	ATOM	771	CE	MET	95	33.754	90.606	12.879	1.00	15.55
	ATOM	772	C	MET	95	32.306	87.074	16.326	1.00	36.36
50	ATOM	773	O	MET	95	31.807	86.765	17.412	1.00	40.70
	ATOM	774	N	PRO	96	33.297	86.359	15.768	1.00	36.06
	ATOM	775	CD	PRO	96	33.684	86.402	14.358	1.00	35.32
	ATOM	776	CA	PRO	96	33.900	85.191	16.411	1.00	40.00
	ATOM	777	CB	PRO	96	34.457	84.385	15.242	1.00	33.97
55	ATOM	778	CG	PRO	96	33.787	84.947	14.053	1.00	37.05
	ATOM	779	C	PRO	96	35.029	85.704	17.302	1.00	46.34
	ATOM	780	O	PRO	96	35.422	86.865	17.197	1.00	47.59
	ATOM	781	N	GLU	97	35.565	84.846	18.164	1.00	52.14
	ATOM	782	CA	GLU	97	36.647	85.256	19.051	1.00	57.27
60	ATOM	783	CB	GLU	97	36.656	84.396	20.315	1.00	62.29
	ATOM	784	CG	GLU	97	37.331	85.046	21.519	1.00	72.31
	ATOM	785	CD	GLU	97	36.315	85.657	22.495	1.00	79.34
	ATOM	786	OE1	GLU	97	35.476	84.886	23.057	1.00	84.94
	ATOM	787	OE2	GLU	97	36.352	86.903	22.696	1.00	81.39
65	ATOM	788	C	GLU	97	37.977	85.089	18.325	1.00	56.67
	ATOM	789	O	GLU	97	38.898	85.893	18.493	1.00	59.18

	ATOM	790	N	THR	98	38.062	84.051	17.501	1.00	57.55
	ATOM	791	CA	THR	98	39.285	83.737	16.771	1.00	59.71
	ATOM	792	CB	THR	98	39.182	82.360	16.128	1.00	59.24
	ATOM	793	OG1	THR	98	38.047	82.354	15.240	1.00	68.81
5	ATOM	794	CG2	THR	98	39.031	81.276	17.215	1.00	56.15
	ATOM	795	C	THR	98	39.724	84.721	15.688	1.00	60.32
	ATOM	796	O	THR	98	40.395	85.715	15.990	1.00	65.03
	ATOM	797	N	LEU	99	39.371	84.439	14.433	1.00	54.94
	ATOM	798	CA	LEU	99	39.768	85.298	13.314	1.00	51.13
10	ATOM	799	CB	LEU	99	40.386	84.444	12.209	1.00	50.56
	ATOM	800	CG	LEU	99	41.783	83.890	12.475	1.00	52.80
	ATOM	801	CD1	LEU	99	42.087	82.738	11.519	1.00	48.38
	ATOM	802	CD2	LEU	99	42.799	85.012	12.321	1.00	50.80
	ATOM	803	C	LEU	99	38.625	86.136	12.733	1.00	50.20
15	ATOM	804	O	LEU	99	38.083	85.816	11.665	1.00	50.50
	ATOM	805	N	PRO	100	38.268	87.243	13.409	1.00	47.97
	ATOM	806	CD	PRO	100	38.954	87.837	14.567	1.00	45.68
	ATOM	807	CA	PRO	100	37.183	88.117	12.955	1.00	46.54
	ATOM	808	CB	PRO	100	37.252	89.296	13.927	1.00	45.50
20	ATOM	809	CG	PRO	100	37.874	88.703	15.149	1.00	45.80
	ATOM	810	C	PRO	100	37.347	88.566	11.513	1.00	46.78
	ATOM	811	O	PRO	100	38.357	89.156	11.149	1.00	52.13
	ATOM	812	N	ASN	101	36.348	88.282	10.695	1.00	42.71
	ATOM	813	CA	ASN	101	36.358	88.678	9.300	1.00	41.37
25	ATOM	814	CB	ASN	101	37.306	87.792	8.501	1.00	43.24
	ATOM	815	CG	ASN	101	38.745	88.261	8.574	1.00	45.93
	ATOM	816	OD1	ASN	101	39.101	89.303	8.020	1.00	45.33
	ATOM	817	ND2	ASN	101	39.585	87.494	9.267	1.00	48.11
	ATOM	818	C	ASN	101	34.946	88.547	8.764	1.00	42.66
30	ATOM	819	O	ASN	101	34.599	87.543	8.139	1.00	44.79
	ATOM	820	N	ASN	102	34.115	89.555	9.011	1.00	40.46
	ATOM	821	CA	ASN	102	32.758	89.470	8.525	1.00	37.90
	ATOM	822	CB	ASN	102	31.772	89.550	9.680	1.00	34.01
	ATOM	823	CG	ASN	102	31.504	88.190	10.294	1.00	32.44
35	ATOM	824	OD1	ASN	102	31.382	87.196	9.583	1.00	29.07
	ATOM	825	ND2	ASN	102	31.405	88.140	11.614	1.00	33.79
	ATOM	826	C	ASN	102	32.316	90.388	7.402	1.00	39.20
	ATOM	827	O	ASN	102	31.995	89.909	6.321	1.00	46.07
	ATOM	828	N	SER	103	32.292	91.690	7.609	1.00	35.42
40	ATOM	829	CA	SER	103	31.821	92.566	6.522	1.00	40.28
	ATOM	830	CB	SER	103	32.582	92.303	5.199	1.00	40.68
	ATOM	831	OG	SER	103	31.829	91.528	4.273	1.00	31.76
	ATOM	832	C	SER	103	30.301	92.385	6.296	1.00	35.24
	ATOM	833	O	SER	103	29.807	91.290	6.034	1.00	24.65
45	ATOM	834	N	CYS	104	29.572	93.484	6.437	1.00	32.88
	ATOM	835	CA	CYS	104	28.142	93.481	6.266	1.00	31.77
	ATOM	836	C	CYS	104	27.783	94.562	5.231	1.00	30.58
	ATOM	837	O	CYS	104	28.311	95.670	5.265	1.00	29.23
	ATOM	838	CB	CYS	104	27.430	93.744	7.622	1.00	28.43
50	ATOM	839	SG	CYS	104	25.701	94.112	7.281	1.00	46.07
	ATOM	840	N	TYR	105	26.907	94.218	4.289	1.00	30.34
	ATOM	841	CA	TYR	105	26.442	95.152	3.265	1.00	26.12
	ATOM	842	CB	TYR	105	26.559	94.529	1.875	1.00	27.03
	ATOM	843	CG	TYR	105	25.925	95.340	0.756	1.00	25.61
55	ATOM	844	CD1	TYR	105	26.691	96.185	-0.054	1.00	24.90
	ATOM	845	CE1	TYR	105	26.112	96.926	-1.074	1.00	26.08
	ATOM	846	CD2	TYR	105	24.558	95.267	0.507	1.00	25.41
	ATOM	847	CE2	TYR	105	23.972	96.009	-0.510	1.00	28.91
	ATOM	848	CZ	TYR	105	24.754	96.835	-1.297	1.00	27.58
60	ATOM	849	OH	TYR	105	24.171	97.559	-2.313	1.00	29.89
	ATOM	850	C	TYR	105	24.975	95.471	3.542	1.00	29.20
	ATOM	851	O	TYR	105	24.216	94.612	3.980	1.00	33.10
	ATOM	852	N	SER	106	24.571	96.705	3.290	1.00	30.48
	ATOM	853	CA	SER	106	23.186	97.086	3.515	1.00	29.22
65	ATOM	854	CB	SER	106	22.971	97.436	4.990	1.00	28.30
	ATOM	855	OG	SER	106	21.622	97.763	5.258	1.00	28.62

	ATOM	856	C	SER	106	22.879	98.284	2.633	1.00	29.94
	ATOM	857	O	SER	106	23.737	99.136	2.438	1.00	29.72
	ATOM	858	N	ALA	107	21.668	98.335	2.085	1.00	27.64
	ATOM	859	CA	ALA	107	21.267	99.440	1.224	1.00	24.33
5	ATOM	860	CB	ALA	107	21.734	99.191	-0.199	1.00	21.30
	ATOM	861	C	ALA	107	19.762	99.655	1.244	1.00	25.56
	ATOM	862	O	ALA	107	19.005	98.810	1.703	1.00	28.03
	ATOM	863	N	GLY	108	19.333	100.800	0.738	1.00	26.07
	ATOM	864	CA	GLY	108	17.918	101.104	0.693	1.00	25.49
10	ATOM	865	C	GLY	108	17.687	102.421	-0.011	1.00	29.03
	ATOM	866	O	GLY	108	18.638	103.095	-0.382	1.00	32.20
	ATOM	867	N	ILE	109	16.428	102.792	-0.197	1.00	29.34
	ATOM	868	CA	ILE	109	16.091	104.041	-0.863	1.00	27.71
	ATOM	869	CB	ILE	109	15.014	103.817	-1.937	1.00	27.48
15	ATOM	870	CG2	ILE	109	14.658	105.128	-2.606	1.00	26.46
	ATOM	871	CG1	ILE	109	15.527	102.815	-2.970	1.00	26.16
	ATOM	872	CD1	ILE	109	14.496	102.392	-3.977	1.00	29.07
	ATOM	873	C	ILE	109	15.576	105.041	0.154	1.00	28.66
	ATOM	874	O	ILE	109	14.874	104.676	1.095	1.00	30.59
20	ATOM	875	N	ALA	110	15.929	106.306	-0.028	1.00	25.60
	ATOM	876	CA	ALA	110	15.491	107.360	0.879	1.00	26.36
	ATOM	877	CB	ALA	110	16.501	107.543	1.987	1.00	20.26
	ATOM	878	C	ALA	110	15.328	108.663	0.117	1.00	29.61
	ATOM	879	O	ALA	110	15.951	108.863	-0.915	1.00	27.61
25	ATOM	880	N	LYS	111	14.473	109.548	0.609	1.00	33.77
	ATOM	881	CA	LYS	111	14.305	110.820	-0.054	1.00	35.09
	ATOM	882	CB	LYS	111	12.857	111.292	-0.005	1.00	39.38
	ATOM	883	CG	LYS	111	12.726	112.704	-0.561	1.00	49.74
	ATOM	884	CD	LYS	111	11.431	112.936	-1.290	1.00	53.95
30	ATOM	885	CE	LYS	111	11.449	114.302	-1.961	1.00	54.28
	ATOM	886	NZ	LYS	111	10.152	114.564	-2.673	1.00	60.85
	ATOM	887	C	LYS	111	15.211	111.834	0.634	1.00	35.28
	ATOM	888	O	LYS	111	15.161	112.000	1.847	1.00	36.58
	ATOM	889	N	LEU	112	16.045	112.498	-0.155	1.00	32.32
35	ATOM	890	CA	LEU	112	16.979	113.479	0.363	1.00	29.67
	ATOM	891	CB	LEU	112	18.410	113.041	0.060	1.00	27.04
	ATOM	892	CG	LEU	112	18.771	111.605	0.436	1.00	24.38
	ATOM	893	CD1	LEU	112	20.169	111.283	-0.051	1.00	27.56
	ATOM	894	CD2	LEU	112	18.669	111.426	1.926	1.00	19.52
40	ATOM	895	C	LEU	112	16.719	114.837	-0.271	1.00	32.58
	ATOM	896	O	LEU	112	16.021	114.937	-1.279	1.00	31.51
	ATOM	897	N	GLU	113	17.289	115.881	0.319	1.00	35.75
	ATOM	898	CA	GLU	113	17.113	117.229	-0.191	1.00	40.08
	ATOM	899	CB	GLU	113	16.214	118.041	0.723	1.00	44.23
45	ATOM	900	CG	GLU	113	14.946	117.368	1.158	1.00	50.12
	ATOM	901	CD	GLU	113	13.925	118.391	1.600	1.00	58.08
	ATOM	902	OE1	GLU	113	14.331	119.403	2.244	1.00	61.13
	ATOM	903	OE2	GLU	113	12.721	118.182	1.300	1.00	63.64
	ATOM	904	C	GLU	113	18.423	117.970	-0.291	1.00	41.63
50	ATOM	905	O	GLU	113	19.399	117.609	0.353	1.00	48.29
	ATOM	906	N	GLU	114	18.423	119.024	-1.101	1.00	40.97
	ATOM	907	CA	GLU	114	19.589	119.879	-1.280	1.00	36.68
	ATOM	908	CB	GLU	114	19.189	121.174	-1.954	1.00	41.19
	ATOM	909	CG	GLU	114	19.377	121.245	-3.427	1.00	46.58
55	ATOM	910	CD	GLU	114	19.565	122.682	-3.863	1.00	51.12
	ATOM	911	OE1	GLU	114	20.606	123.276	-3.489	1.00	51.88
	ATOM	912	OE2	GLU	114	18.670	123.218	-4.561	1.00	57.08
	ATOM	913	C	GLU	114	20.127	120.259	0.084	1.00	35.51
	ATOM	914	O	GLU	114	19.405	120.831	0.892	1.00	38.92
60	ATOM	915	N	GLY	115	21.391	119.963	0.340	1.00	32.99
	ATOM	916	CA	GLY	115	21.952	120.334	1.620	1.00	31.79
	ATOM	917	C	GLY	115	22.049	119.197	2.601	1.00	34.39
	ATOM	918	O	GLY	115	22.768	119.303	3.585	1.00	39.20
	ATOM	919	N	ASP	116	21.311	118.122	2.358	1.00	35.66
65	ATOM	920	CA	ASP	116	21.375	116.971	3.242	1.00	38.09
	ATOM	921	CB	ASP	116	20.323	115.920	2.867	1.00	36.00

	ATOM	922	CG	ASP	116	18.929	116.287	3.326	1.00	35.09
	ATOM	923	OD1	ASP	116	18.785	117.188	4.178	1.00	33.81
	ATOM	924	OD2	ASP	116	17.971	115.655	2.844	1.00	35.24
	ATOM	925	C	ASP	116	22.758	116.356	3.088	1.00	39.38
5	ATOM	926	O	ASP	116	23.396	116.492	2.040	1.00	37.91
	ATOM	927	N	GLU	117	23.231	115.692	4.135	1.00	41.54
	ATOM	928	CA	GLU	117	24.528	115.032	4.073	1.00	37.92
	ATOM	929	CB	GLU	117	25.531	115.711	4.998	1.00	38.90
	ATOM	930	CG	GLU	117	25.746	117.173	4.702	1.00	48.15
10	ATOM	931	CD	GLU	117	26.780	117.794	5.618	1.00	52.62
	ATOM	932	OE1	GLU	117	26.814	117.428	6.825	1.00	55.49
	ATOM	933	OE2	GLU	117	27.551	118.655	5.128	1.00	57.10
	ATOM	934	C	GLU	117	24.342	113.591	4.506	1.00	34.02
	ATOM	935	O	GLU	117	23.522	113.303	5.376	1.00	33.31
15	ATOM	936	N	LEU	118	25.081	112.687	3.878	1.00	30.68
	ATOM	937	CA	LEU	118	25.020	111.277	4.224	1.00	27.14
	ATOM	938	CB	LEU	118	24.794	110.426	2.974	1.00	26.49
	ATOM	939	CG	LEU	118	23.473	110.570	2.227	1.00	26.21
	ATOM	940	CD1	LEU	118	23.503	109.707	0.987	1.00	23.05
20	ATOM	941	CD2	LEU	118	22.323	110.169	3.127	1.00	25.38
	ATOM	942	C	LEU	118	26.349	110.882	4.855	1.00	28.33
	ATOM	943	O	LEU	118	27.400	111.399	4.472	1.00	31.46
	ATOM	944	N	GLN	119	26.311	109.980	5.830	1.00	28.96
	ATOM	945	CA	GLN	119	27.542	109.516	6.457	1.00	29.13
25	ATOM	946	CB	GLN	119	27.918	110.424	7.619	1.00	28.23
	ATOM	947	CG	GLN	119	27.005	110.315	8.802	1.00	36.38
	ATOM	948	CD	GLN	119	27.376	111.281	9.901	1.00	36.57
	ATOM	949	OE1	GLN	119	26.928	111.136	11.038	1.00	40.70
	ATOM	950	NE2	GLN	119	28.188	112.278	9.571	1.00	36.80
30	ATOM	951	C	GLN	119	27.433	108.070	6.930	1.00	29.43
	ATOM	952	O	GLN	119	26.343	107.578	7.212	1.00	28.80
	ATOM	953	N	LEU	120	28.578	107.399	6.990	1.00	32.26
	ATOM	954	CA	LEU	120	28.680	106.006	7.416	1.00	29.96
	ATOM	955	CB	LEU	120	29.532	105.224	6.405	1.00	31.17
35	ATOM	956	CG	LEU	120	29.653	103.696	6.349	1.00	30.26
	ATOM	957	CD1	LEU	120	29.900	103.116	7.719	1.00	27.10
	ATOM	958	CD2	LEU	120	28.397	103.133	5.747	1.00	30.96
	ATOM	959	C	LEU	120	29.367	106.014	8.783	1.00	30.51
	ATOM	960	O	LEU	120	30.540	106.372	8.899	1.00	32.88
40	ATOM	961	N	ALA	121	28.635	105.613	9.816	1.00	27.40
	ATOM	962	CA	ALA	121	29.175	105.600	11.169	1.00	25.55
	ATOM	963	CB	ALA	121	28.393	106.577	12.032	1.00	19.41
	ATOM	964	C	ALA	121	29.184	104.222	11.832	1.00	29.89
	ATOM	965	O	ALA	121	28.234	103.452	11.707	1.00	35.88
45	ATOM	966	N	ILE	122	30.274	103.921	12.531	1.00	30.33
	ATOM	967	CA	ILE	122	30.408	102.660	13.248	1.00	30.27
	ATOM	968	CB	ILE	122	31.762	101.997	12.977	1.00	29.27
	ATOM	969	CG2	ILE	122	31.811	100.654	13.668	1.00	30.38
	ATOM	970	CG1	ILE	122	31.965	101.824	11.469	1.00	26.03
50	ATOM	971	CD1	ILE	122	33.293	101.204	11.075	1.00	29.21
	ATOM	972	C	ILE	122	30.306	103.010	14.729	1.00	33.20
	ATOM	973	O	ILE	122	31.164	103.707	15.266	1.00	33.30
	ATOM	974	N	PRO	123	29.246	102.534	15.406	1.00	34.70
	ATOM	975	CD	PRO	123	28.161	101.733	14.831	1.00	34.40
55	ATOM	976	CA	PRO	123	28.990	102.783	16.831	1.00	36.49
	ATOM	977	CB	PRO	123	27.538	102.331	17.015	1.00	30.22
	ATOM	978	CG	PRO	123	26.994	102.227	15.619	1.00	36.72
	ATOM	979	C	PRO	123	29.920	102.027	17.778	1.00	39.64
	ATOM	980	O	PRO	123	29.451	101.274	18.637	1.00	40.48
60	ATOM	981	N	ARG	124	31.226	102.227	17.624	1.00	44.58
	ATOM	982	CA	ARG	124	32.206	101.560	18.469	1.00	48.04
	ATOM	983	CB	ARG	124	32.600	100.217	17.865	1.00	52.91
	ATOM	984	CG	ARG	124	31.422	99.303	17.582	1.00	60.50
	ATOM	985	CD	ARG	124	31.440	98.158	18.571	1.00	72.57
65	ATOM	986	NE	ARG	124	30.235	97.317	18.465	1.00	84.90
	ATOM	987	CZ	ARG	124	29.105	97.491	19.162	1.00	85.44

	ATOM	988	NH1	ARG	124	29.005	98.493	20.037	1.00	87.69
	ATOM	989	NH2	ARG	124	28.070	96.661	18.979	1.00	82.78
	ATOM	990	C	ARG	124	33.446	102.414	18.677	1.00	48.39
	ATOM	991	O	ARG	124	33.839	103.175	17.792	1.00	45.26
5	ATOM	992	N	GLU	125	34.051	102.264	19.853	1.00	52.84
	ATOM	993	CA	GLU	125	35.240	103.014	20.250	1.00	55.02
	ATOM	994	CB	GLU	125	35.770	102.468	21.579	1.00	63.62
	ATOM	995	CG	GLU	125	35.479	100.982	21.803	1.00	70.62
	ATOM	996	CD	GLU	125	33.996	100.651	21.685	1.00	74.76
10	ATOM	997	OE1	GLU	125	33.202	101.139	22.534	1.00	75.75
	ATOM	998	OE2	GLU	125	33.628	99.906	20.743	1.00	80.71
	ATOM	999	C	GLU	125	36.362	103.047	19.221	1.00	53.44
	ATOM	1000	O	GLU	125	36.631	104.096	18.635	1.00	57.50
	ATOM	1001	N	ASN	126	37.050	101.931	19.025	1.00	47.18
15	ATOM	1002	CA	ASN	126	38.109	101.903	18.025	1.00	47.59
	ATOM	1003	CB	ASN	126	39.494	101.864	18.665	1.00	51.03
	ATOM	1004	CG	ASN	126	40.174	103.225	18.654	1.00	57.07
	ATOM	1005	OD1	ASN	126	39.838	104.120	19.445	1.00	61.15
	ATOM	1006	ND2	ASN	126	41.127	103.397	17.737	1.00	56.63
20	ATOM	1007	C	ASN	126	37.896	100.686	17.167	1.00	47.06
	ATOM	1008	O	ASN	126	38.653	99.713	17.221	1.00	45.61
	ATOM	1009	N	ALA	127	36.833	100.758	16.377	1.00	43.50
	ATOM	1010	CA	ALA	127	36.444	99.675	15.502	1.00	38.26
	ATOM	1011	CB	ALA	127	35.382	100.160	14.543	1.00	37.97
25	ATOM	1012	C	ALA	127	37.617	99.103	14.731	1.00	37.16
	ATOM	1013	O	ALA	127	38.376	99.833	14.096	1.00	33.14
	ATOM	1014	N	GLN	128	37.773	97.787	14.814	1.00	37.87
	ATOM	1015	CA	GLN	128	38.823	97.106	14.071	1.00	37.19
	ATOM	1016	CB	GLN	128	39.229	95.830	14.801	1.00	40.23
30	ATOM	1017	CG	GLN	128	39.939	96.122	16.102	1.00	38.14
	ATOM	1018	CD	GLN	128	41.118	97.041	15.886	1.00	39.51
	ATOM	1019	OE1	GLN	128	42.115	96.660	15.263	1.00	40.36
	ATOM	1020	NE2	GLN	128	41.006	98.274	16.379	1.00	41.53
	ATOM	1021	C	GLN	128	38.199	96.809	12.710	1.00	35.55
35	ATOM	1022	O	GLN	128	37.430	95.859	12.538	1.00	35.57
	ATOM	1023	N	ILE	129	38.543	97.656	11.752	1.00	34.64
	ATOM	1024	CA	ILE	129	37.994	97.611	10.404	1.00	29.60
	ATOM	1025	CB	ILE	129	37.161	98.927	10.192	1.00	31.03
	ATOM	1026	CG2	ILE	129	37.471	99.597	8.878	1.00	29.76
40	ATOM	1027	CG1	ILE	129	35.689	98.619	10.350	1.00	26.37
	ATOM	1028	CD1	ILE	129	35.354	98.103	11.699	1.00	34.70
	ATOM	1029	C	ILE	129	39.065	97.470	9.327	1.00	29.55
	ATOM	1030	O	ILE	129	40.209	97.848	9.531	1.00	33.42
	ATOM	1031	N	SER	130	38.696	96.913	8.179	1.00	27.67
45	ATOM	1032	CA	SER	130	39.638	96.791	7.071	1.00	25.00
	ATOM	1033	CB	SER	130	39.356	95.549	6.243	1.00	22.09
	ATOM	1034	OG	SER	130	40.124	95.579	5.048	1.00	20.89
	ATOM	1035	C	SER	130	39.468	98.014	6.184	1.00	26.03
	ATOM	1036	O	SER	130	38.351	98.410	5.901	1.00	30.77
50	ATOM	1037	N	LEU	131	40.564	98.609	5.738	1.00	24.49
	ATOM	1038	CA	LEU	131	40.475	99.794	4.899	1.00	28.07
	ATOM	1039	CB	LEU	131	41.363	100.906	5.458	1.00	28.87
	ATOM	1040	CG	LEU	131	40.726	101.831	6.494	1.00	30.76
	ATOM	1041	CD1	LEU	131	40.005	101.046	7.550	1.00	28.98
55	ATOM	1042	CD2	LEU	131	41.805	102.681	7.112	1.00	33.37
	ATOM	1043	C	LEU	131	40.810	99.564	3.435	1.00	33.66
	ATOM	1044	O	LEU	131	41.312	100.464	2.761	1.00	37.67
	ATOM	1045	N	ASP	132	40.535	98.361	2.943	1.00	34.97
	ATOM	1046	CA	ASP	132	40.784	98.042	1.541	1.00	34.31
60	ATOM	1047	CB	ASP	132	40.955	96.536	1.346	1.00	42.68
	ATOM	1048	CG	ASP	132	42.314	96.036	1.807	1.00	47.98
	ATOM	1049	OD1	ASP	132	42.494	94.796	1.893	1.00	50.41
	ATOM	1050	OD2	ASP	132	43.201	96.886	2.071	1.00	46.42
	ATOM	1051	C	ASP	132	39.607	98.543	0.721	1.00	34.32
65	ATOM	1052	O	ASP	132	38.454	98.322	1.074	1.00	32.87
	ATOM	1053	N	GLY	133	39.914	99.222	-0.376	1.00	34.53

	ATOM	1054	CA	GLY	133	38.879	99.778	-1.224	1.00	36.93
	ATOM	1055	C	GLY	133	37.809	98.830	-1.728	1.00	35.06
	ATOM	1056	O	GLY	133	36.716	99.266	-2.089	1.00	40.44
	ATOM	1057	N	ASP	134	38.100	97.538	-1.754	1.00	31.08
5	ATOM	1058	CA	ASP	134	37.123	96.590	-2.252	1.00	27.63
	ATOM	1059	CB	ASP	134	37.822	95.487	-3.051	1.00	28.43
	ATOM	1060	CG	ASP	134	38.824	94.704	-2.231	1.00	31.78
	ATOM	1061	OD1	ASP	134	39.462	95.295	-1.340	1.00	35.68
	ATOM	1062	OD2	ASP	134	38.989	93.494	-2.491	1.00	32.53
10	ATOM	1063	C	ASP	134	36.220	96.004	-1.182	1.00	25.08
	ATOM	1064	O	ASP	134	35.144	95.538	-1.483	1.00	19.71
	ATOM	1065	N	VAL	135	36.627	96.058	0.076	1.00	26.84
	ATOM	1066	CA	VAL	135	35.781	95.500	1.121	1.00	23.72
	ATOM	1067	CB	VAL	135	36.596	94.666	2.112	1.00	25.13
15	ATOM	1068	CG1	VAL	135	37.054	93.405	1.432	1.00	22.76
	ATOM	1069	CG2	VAL	135	37.778	95.461	2.623	1.00	22.12
	ATOM	1070	C	VAL	135	34.928	96.509	1.883	1.00	26.54
	ATOM	1071	O	VAL	135	33.895	96.142	2.427	1.00	29.26
	ATOM	1072	N	THR	136	35.346	97.772	1.942	1.00	26.39
20	ATOM	1073	CA	THR	136	34.521	98.770	2.621	1.00	28.40
	ATOM	1074	CB	THR	136	35.072	99.121	4.044	1.00	28.21
	ATOM	1075	OG1	THR	136	35.743	100.379	4.017	1.00	37.13
	ATOM	1076	CG2	THR	136	36.024	98.056	4.527	1.00	26.90
	ATOM	1077	C	THR	136	34.358	100.022	1.747	1.00	31.60
25	ATOM	1078	O	THR	136	35.329	100.709	1.417	1.00	27.29
	ATOM	1079	N	PHE	137	33.110	100.289	1.359	1.00	31.45
	ATOM	1080	CA	PHE	137	32.785	101.412	0.488	1.00	26.64
	ATOM	1081	CB	PHE	137	32.798	100.928	-0.955	1.00	26.84
	ATOM	1082	CG	PHE	137	32.135	99.596	-1.152	1.00	24.32
30	ATOM	1083	CD1	PHE	137	30.762	99.508	-1.358	1.00	27.77
	ATOM	1084	CD2	PHE	137	32.881	98.429	-1.142	1.00	23.70
	ATOM	1085	CE1	PHE	137	30.146	98.283	-1.553	1.00	27.32
	ATOM	1086	CE2	PHE	137	32.271	97.199	-1.335	1.00	26.31
	ATOM	1087	CZ	PHE	137	30.901	97.129	-1.542	1.00	29.43
35	ATOM	1088	C	PHE	137	31.449	102.061	0.842	1.00	28.06
	ATOM	1089	O	PHE	137	30.660	101.486	1.577	1.00	25.04
	ATOM	1090	N	PHE	138	31.202	103.257	0.308	1.00	29.81
	ATOM	1091	CA	PHE	138	29.990	104.009	0.623	1.00	29.93
	ATOM	1092	CB	PHE	138	30.388	105.165	1.539	1.00	27.47
40	ATOM	1093	CG	PHE	138	29.240	105.861	2.187	1.00	28.49
	ATOM	1094	CD1	PHE	138	28.019	105.220	2.364	1.00	27.65
	ATOM	1095	CD2	PHE	138	29.384	107.173	2.632	1.00	31.06
	ATOM	1096	CE1	PHE	138	26.958	105.878	2.972	1.00	30.90
	ATOM	1097	CE2	PHE	138	28.331	107.843	3.242	1.00	27.65
45	ATOM	1098	CZ	PHE	138	27.115	107.196	3.412	1.00	27.41
	ATOM	1099	C	PHE	138	29.211	104.493	-0.616	1.00	30.91
	ATOM	1100	O	PHE	138	29.757	105.132	-1.508	1.00	24.94
	ATOM	1101	N	GLY	139	27.911	104.185	-0.604	1.00	36.94
	ATOM	1102	CA	GLY	139	26.960	104.435	-1.686	1.00	36.08
50	ATOM	1103	C	GLY	139	26.512	105.731	-2.315	1.00	36.48
	ATOM	1104	O	GLY	139	27.334	106.546	-2.691	1.00	43.61
	ATOM	1105	N	ALA	140	25.196	105.865	-2.490	1.00	36.60
	ATOM	1106	CA	ALA	140	24.533	107.038	-3.092	1.00	34.94
	ATOM	1107	CB	ALA	140	25.158	108.320	-2.572	1.00	36.94
55	ATOM	1108	C	ALA	140	24.428	107.108	-4.628	1.00	32.53
	ATOM	1109	O	ALA	140	25.392	107.405	-5.325	1.00	27.96
	ATOM	1110	N	LEU	141	23.223	106.855	-5.133	1.00	33.23
	ATOM	1111	CA	LEU	141	22.913	106.889	-6.569	1.00	36.11
	ATOM	1112	CB	LEU	141	22.891	105.466	-7.126	1.00	37.53
60	ATOM	1113	CG	LEU	141	22.505	105.192	-8.583	1.00	38.87
	ATOM	1114	CD1	LEU	141	22.826	103.753	-8.906	1.00	40.67
	ATOM	1115	CD2	LEU	141	21.030	105.445	-8.824	1.00	37.53
	ATOM	1116	C	LEU	141	21.536	107.536	-6.749	1.00	38.89
	ATOM	1117	O	LEU	141	20.571	107.133	-6.103	1.00	39.25
65	ATOM	1118	N	LYS	142	21.426	108.529	-7.625	1.00	42.86
	ATOM	1119	CA	LYS	142	20.134	109.182	-7.809	1.00	42.96

385

	ATOM	1120	CB	LYS	142	20.329	110.626	-8.260	1.00	41.80
	ATOM	1121	CG	LYS	142	19.015	111.334	-8.491	1.00	46.88
	ATOM	1122	CD	LYS	142	19.151	112.842	-8.470	1.00	49.58
	ATOM	1123	CE	LYS	142	17.791	113.478	-8.705	1.00	51.07
5	ATOM	1124	NZ	LYS	142	17.833	114.958	-8.615	1.00	56.20
	ATOM	1125	C	LYS	142	19.183	108.473	-8.767	1.00	43.41
	ATOM	1126	O	LYS	142	19.551	108.137	-9.889	1.00	44.98
	ATOM	1127	N	LEU	143	17.952	108.253	-8.315	1.00	42.81
	ATOM	1128	CA	LEU	143	16.937	107.589	-9.130	1.00	40.65
10	ATOM	1129	CB	LEU	143	15.890	106.926	-8.237	1.00	36.56
	ATOM	1130	CG	LEU	143	16.358	105.896	-7.209	1.00	36.17
	ATOM	1131	CD1	LEU	143	15.164	105.422	-6.401	1.00	37.95
	ATOM	1132	CD2	LEU	143	17.025	104.731	-7.907	1.00	27.38
	ATOM	1133	C	LEU	143	16.234	108.597	-10.032	1.00	43.14
15	ATOM	1134	O	LEU	143	16.086	109.769	-9.670	1.00	46.28
	ATOM	1135	N	LEU	144	15.799	108.149	-11.202	1.00	42.63
	ATOM	1136	CA	LEU	144	15.100	109.037	-12.119	1.00	44.11
	ATOM	1137	CB	LEU	144	15.133	108.478	-13.538	1.00	44.88
	ATOM	1138	CG	LEU	144	16.528	108.405	-14.156	1.00	47.02
20	ATOM	1139	CD1	LEU	144	16.458	107.749	-15.508	1.00	47.71
	ATOM	1140	CD2	LEU	144	17.109	109.802	-14.276	1.00	47.87
	ATOM	1141	C	LEU	144	13.658	109.198	-11.680	1.00	45.66
	ATOM	1142	O	LEU	144	13.168	108.333	-10.931	1.00	46.37
	ATOM	1143	OXT	LEU	144	13.031	110.187	-12.104	1.00	50.66
25	END					-16.719	146.167	89.779	0.00	0.00

TABLE 8

	111								
	ATOM	1	CB	VAL	1	9.561	110.300	-17.231	1.00 69.28
5	ATOM	2	CG1	VAL	1	10.302	111.648	-17.092	1.00 68.26
	ATOM	3	CG2	VAL	1	8.686	110.275	-18.489	1.00 71.75
	ATOM	4	C	VAL	1	11.152	108.942	-15.841	1.00 64.86
	ATOM	5	O	VAL	1	12.297	109.330	-15.564	1.00 64.02
	ATOM	6	N	VAL	1	9.948	107.858	-17.745	1.00 65.10
10	ATOM	7	CA	VAL	1	10.586	109.122	-17.263	1.00 65.98
	ATOM	8	N	THR	2	10.347	108.368	-14.944	1.00 61.42
	ATOM	9	CA	THR	2	10.781	108.144	-13.566	1.00 56.64
	ATOM	10	CB	THR	2	9.883	108.875	-12.553	1.00 56.77
	ATOM	11	OG1	THR	2	8.525	108.477	-12.761	1.00 57.45
15	ATOM	12	CG2	THR	2	10.001	110.386	-12.706	1.00 55.68
	ATOM	13	C	THR	2	10.753	106.672	-13.202	1.00 53.08
	ATOM	14	O	THR	2	10.214	105.848	-13.934	1.00 52.00
	ATOM	15	N	GLN	3	11.325	106.351	-12.049	1.00 50.34
	ATOM	16	CA	GLN	3	11.368	104.974	-11.574	1.00 45.97
20	ATOM	17	CB	GLN	3	12.800	104.585	-11.228	1.00 46.09
	ATOM	18	CG	GLN	3	13.802	104.928	-12.299	1.00 49.60
	ATOM	19	CD	GLN	3	15.189	104.453	-11.947	1.00 50.43
	ATOM	20	OE1	GLN	3	15.441	103.254	-11.898	1.00 50.94
	ATOM	21	NE2	GLN	3	16.097	105.392	-11.692	1.00 49.99
25	ATOM	22	C	GLN	3	10.492	104.795	-10.348	1.00 42.41
	ATOM	23	O	GLN	3	10.804	105.287	-9.267	1.00 37.34
	ATOM	24	N	ASP	4	9.388	104.087	-10.521	1.00 40.32
	ATOM	25	CA	ASP	4	8.477	103.848	-9.417	1.00 39.37
	ATOM	26	CB	ASP	4	7.187	103.201	-9.929	1.00 44.56
30	ATOM	27	CG	ASP	4	6.395	104.119	-10.835	1.00 45.76
	ATOM	28	OD1	ASP	4	6.992	105.089	-11.356	1.00 48.58
	ATOM	29	OD2	ASP	4	5.187	103.869	-11.031	1.00 45.55
	ATOM	30	C	ASP	4	9.142	102.929	-8.416	1.00 36.30
	ATOM	31	O	ASP	4	9.931	102.062	-8.783	1.00 36.29
35	ATOM	32	N	CYS	5	8.830	103.132	-7.146	1.00 35.11
	ATOM	33	CA	CYS	5	9.373	102.294	-6.092	1.00 35.61
	ATOM	34	CB	CYS	5	10.827	102.662	-5.787	1.00 34.96
	ATOM	35	SG	CYS	5	11.175	104.408	-5.727	1.00 35.34
	ATOM	36	C	CYS	5	8.514	102.417	-4.851	1.00 31.80
40	ATOM	37	O	CYS	5	7.870	103.435	-4.628	1.00 31.85
	ATOM	38	N	LEU	6	8.480	101.352	-4.067	1.00 27.72
	ATOM	39	CA	LEU	6	7.711	101.323	-2.840	1.00 30.80
	ATOM	40	CB	LEU	6	6.402	100.572	-3.054	1.00 29.25
	ATOM	41	CG	LEU	6	5.489	100.374	-1.843	1.00 29.47
45	ATOM	42	CD1	LEU	6	4.067	100.199	-2.304	1.00 30.65
	ATOM	43	CD2	LEU	6	5.926	99.167	-1.069	1.00 35.27
	ATOM	44	C	LEU	6	8.559	100.624	-1.797	1.00 31.83
	ATOM	45	O	LEU	6	9.055	99.529	-2.036	1.00 33.45
	ATOM	46	N	GLN	7	8.739	101.258	-0.644	1.00 32.23
50	ATOM	47	CA	GLN	7	9.547	100.677	0.407	1.00 30.26
	ATOM	48	CB	GLN	7	10.732	101.587	0.705	1.00 29.56
	ATOM	49	CG	GLN	7	11.811	100.969	1.567	1.00 25.09
	ATOM	50	CD	GLN	7	13.058	101.828	1.605	1.00 26.50
	ATOM	51	OE1	GLN	7	13.076	102.902	2.207	1.00 24.40
55	ATOM	52	NE2	GLN	7	14.105	101.367	0.945	1.00 27.03
	ATOM	53	C	GLN	7	8.715	100.465	1.652	1.00 32.11
	ATOM	54	O	GLN	7	7.875	101.286	1.999	1.00 34.24
	ATOM	55	N	LEU	8	8.951	99.339	2.310	1.00 33.73
	ATOM	56	CA	LEU	8	8.242	98.979	3.527	1.00 35.41
60	ATOM	57	CB	LEU	8	7.540	97.622	3.345	1.00 32.44
	ATOM	58	CG	LEU	8	6.080	97.551	2.874	1.00 27.69
	ATOM	59	CD1	LEU	8	5.641	98.834	2.217	1.00 26.43
	ATOM	60	CD2	LEU	8	5.937	96.378	1.944	1.00 18.64
	ATOM	61	C	LEU	8	9.211	98.919	4.698	1.00 36.87
65	ATOM	62	O	LEU	8	10.379	98.563	4.546	1.00 36.05
	ATOM	63	N	ILE	9	8.705	99.263	5.872	1.00 39.25

	ATOM	64	CA	ILE	9	9.493	99.285	7.093	1.00	35.57
	ATOM	65	CB	ILE	9	9.647	100.756	7.557	1.00	34.28
	ATOM	66	CG2	ILE	9	9.387	100.906	9.035	1.00	38.20
	ATOM	67	CG1	ILE	9	11.028	101.256	7.199	1.00	35.22
5	ATOM	68	CD1	ILE	9	11.247	102.650	7.680	1.00	43.63
	ATOM	69	C	ILE	9	8.803	98.433	8.163	1.00	35.28
	ATOM	70	O	ILE	9	7.579	98.352	8.202	1.00	33.75
	ATOM	71	N	ALA	10	9.583	97.784	9.019	1.00	35.78
	ATOM	72	CA	ALA	10	8.995	96.965	10.068	1.00	35.28
10	ATOM	73	CB	ALA	10	10.076	96.242	10.838	1.00	28.47
	ATOM	74	C	ALA	10	8.171	97.825	11.017	1.00	36.49
	ATOM	75	O	ALA	10	8.565	98.931	11.375	1.00	37.41
	ATOM	76	N	ASP	11	7.019	97.309	11.420	1.00	39.44
	ATOM	77	CA	ASP	11	6.138	98.024	12.337	1.00	40.77
15	ATOM	78	CB	ASP	11	4.697	97.968	11.826	1.00	38.90
	ATOM	79	CG	ASP	11	3.718	98.579	12.799	1.00	40.34
	ATOM	80	OD1	ASP	11	4.129	99.453	13.600	1.00	39.50
	ATOM	81	OD2	ASP	11	2.536	98.189	12.753	1.00	37.38
	ATOM	82	C	ASP	11	6.228	97.415	13.731	1.00	42.25
20	ATOM	83	O	ASP	11	5.530	96.452	14.055	1.00	40.54
	ATOM	84	N	SER	12	7.094	97.995	14.553	1.00	42.92
	ATOM	85	CA	SER	12	7.335	97.514	15.912	1.00	43.84
	ATOM	86	CB	SER	12	8.480	98.304	16.536	1.00	42.13
	ATOM	87	OG	SER	12	8.179	99.688	16.547	1.00	43.31
25	ATOM	88	C	SER	12	6.135	97.572	16.842	1.00	46.10
	ATOM	89	O	SER	12	6.205	97.109	17.980	1.00	43.90
	ATOM	90	N	GLU	13	5.033	98.133	16.359	1.00	46.72
	ATOM	91	CA	GLU	13	3.844	98.246	17.182	1.00	44.93
	ATOM	92	CB	GLU	13	3.264	99.643	17.059	1.00	46.71
30	ATOM	93	CG	GLU	13	4.177	100.679	17.657	1.00	58.25
	ATOM	94	CD	GLU	13	3.427	101.922	18.088	1.00	66.70
	ATOM	95	OE1	GLU	13	2.495	101.780	18.928	1.00	74.10
	ATOM	96	OE2	GLU	13	3.761	103.034	17.597	1.00	71.55
	ATOM	97	C	GLU	13	2.762	97.197	16.949	1.00	43.24
35	ATOM	98	O	GLU	13	1.632	97.349	17.417	1.00	44.25
	ATOM	99	N	THR	14	3.093	96.141	16.220	1.00	37.06
	ATOM	100	CA	THR	14	2.138	95.059	16.018	1.00	36.77
	ATOM	101	CB	THR	14	1.363	95.172	14.671	1.00	33.39
	ATOM	102	OG1	THR	14	2.214	94.808	13.586	1.00	41.69
40	ATOM	103	CG2	THR	14	0.858	96.590	14.462	1.00	31.66
	ATOM	104	C	THR	14	2.940	93.763	16.065	1.00	34.96
	ATOM	105	O	THR	14	4.115	93.739	15.719	1.00	35.04
	ATOM	106	N	PRO	15	2.317	92.675	16.516	1.00	37.11
	ATOM	107	CD	PRO	15	0.924	92.603	16.983	1.00	36.50
45	ATOM	108	CA	PRO	15	2.972	91.366	16.623	1.00	38.81
	ATOM	109	CB	PRO	15	1.907	90.499	17.294	1.00	39.15
	ATOM	110	CG	PRO	15	1.006	91.496	17.982	1.00	37.32
	ATOM	111	C	PRO	15	3.378	90.793	15.271	1.00	37.78
	ATOM	112	O	PRO	15	2.748	91.089	14.260	1.00	35.91
50	ATOM	113	N	THR	16	4.422	89.971	15.246	1.00	36.67
	ATOM	114	CA	THR	16	4.835	89.372	13.988	1.00	38.46
	ATOM	115	CB	THR	16	6.161	88.598	14.120	1.00	38.61
	ATOM	116	OG1	THR	16	5.989	87.499	15.021	1.00	42.44
	ATOM	117	CG2	THR	16	7.257	89.513	14.644	1.00	40.11
55	ATOM	118	C	THR	16	3.740	88.394	13.620	1.00	35.97
	ATOM	119	O	THR	16	3.236	87.688	14.483	1.00	39.85
	ATOM	120	N	ILE	17	3.367	88.355	12.347	1.00	33.66
	ATOM	121	CA	ILE	17	2.314	87.457	11.898	1.00	34.76
	ATOM	122	CB	ILE	17	1.771	87.897	10.533	1.00	32.44
60	ATOM	123	CG2	ILE	17	0.763	86.888	10.022	1.00	28.66
	ATOM	124	CG1	ILE	17	1.150	89.294	10.657	1.00	33.15
	ATOM	125	CD1	ILE	17	0.646	89.875	9.352	1.00	31.19
	ATOM	126	C	ILE	17	2.757	86.003	11.802	1.00	40.64
	ATOM	127	O	ILE	17	3.758	85.691	11.171	1.00	44.07
65	ATOM	128	N	GLN	18	2.003	85.109	12.436	1.00	45.91
	ATOM	129	CA	GLN	18	2.315	83.682	12.407	1.00	50.22

	ATOM	130	CB	GLN	18	2.206	83.076	13.802	1.00	53.09
	ATOM	131	CG	GLN	18	3.481	82.410	14.248	1.00	57.60
	ATOM	132	CD	GLN	18	4.569	83.426	14.480	1.00	61.39
	ATOM	133	OE1	GLN	18	4.502	84.218	15.424	1.00	68.36
5	ATOM	134	NE2	GLN	18	5.570	83.429	13.611	1.00	64.93
	ATOM	135	C	GLN	18	1.363	82.945	11.483	1.00	51.56
	ATOM	136	O	GLN	18	0.151	83.129	11.551	1.00	56.44
	ATOM	137	N	LYS	19	1.908	82.101	10.623	1.00	52.60
	ATOM	138	CA	LYS	19	1.066	81.353	9.705	1.00	56.10
10	ATOM	139	CB	LYS	19	0.451	82.284	8.660	1.00	55.55
	ATOM	140	CG	LYS	19	-0.341	81.522	7.623	1.00	61.15
	ATOM	141	CD	LYS	19	-1.143	82.421	6.705	1.00	64.42
	ATOM	142	CE	LYS	19	-1.933	81.575	5.696	1.00	64.84
	ATOM	143	NZ	LYS	19	-2.778	82.405	4.778	1.00	69.52
15	ATOM	144	C	LYS	19	1.818	80.233	9.001	1.00	57.41
	ATOM	145	O	LYS	19	2.934	80.434	8.518	1.00	61.02
	ATOM	146	N	GLY	20	1.192	79.057	8.943	1.00	58.84
	ATOM	147	CA	GLY	20	1.799	77.909	8.297	1.00	56.66
	ATOM	148	C	GLY	20	3.228	77.665	8.740	1.00	56.87
20	ATOM	149	O	GLY	20	4.067	77.283	7.923	1.00	58.64
	ATOM	150	N	SER	21	3.501	77.881	10.028	1.00	54.49
	ATOM	151	CA	SER	21	4.836	77.692	10.599	1.00	53.68
	ATOM	152	CB	SER	21	5.279	76.230	10.452	1.00	55.80
	ATOM	153	OG	SER	21	5.866	75.982	9.188	1.00	65.34
25	ATOM	154	C	SER	21	5.890	78.639	9.976	1.00	51.41
	ATOM	155	O	SER	21	7.096	78.337	9.939	1.00	47.74
	ATOM	156	N	TYR	22	5.412	79.782	9.486	1.00	46.25
	ATOM	157	CA	TYR	22	6.257	80.809	8.894	1.00	40.57
	ATOM	158	CB	TYR	22	5.878	81.042	7.429	1.00	43.27
30	ATOM	159	CG	TYR	22	6.583	80.132	6.456	1.00	49.18
	ATOM	160	CD1	TYR	22	6.450	78.751	6.541	1.00	52.61
	ATOM	161	CE1	TYR	22	7.112	77.903	5.649	1.00	51.92
	ATOM	162	CD2	TYR	22	7.394	80.650	5.453	1.00	50.01
	ATOM	163	CE2	TYR	22	8.058	79.813	4.555	1.00	50.94
35	ATOM	164	CZ	TYR	22	7.913	78.441	4.660	1.00	50.78
	ATOM	165	OH	TYR	22	8.570	77.614	3.776	1.00	53.87
	ATOM	166	C	TYR	22	6.035	82.098	9.678	1.00	37.85
	ATOM	167	O	TYR	22	4.946	82.328	10.202	1.00	38.70
	ATOM	168	N	THR	23	7.063	82.931	9.778	1.00	33.81
40	ATOM	169	CA	THR	23	6.922	84.200	10.484	1.00	32.40
	ATOM	170	CB	THR	23	8.079	84.462	11.481	1.00	31.58
	ATOM	171	OG1	THR	23	8.170	83.389	12.423	1.00	30.76
	ATOM	172	CG2	THR	23	7.837	85.756	12.228	1.00	24.08
	ATOM	173	C	THR	23	6.940	85.309	9.447	1.00	33.28
45	ATOM	174	O	THR	23	7.802	85.331	8.575	1.00	30.44
	ATOM	175	N	PHE	24	5.987	86.223	9.542	1.00	32.16
	ATOM	176	CA	PHE	24	5.914	87.335	8.610	1.00	30.68
	ATOM	177	CB	PHE	24	4.594	87.304	7.840	1.00	28.20
	ATOM	178	CG	PHE	24	4.460	86.128	6.926	1.00	31.00
50	ATOM	179	CD1	PHE	24	4.009	84.908	7.404	1.00	31.69
	ATOM	180	CD2	PHE	24	4.815	86.230	5.584	1.00	31.42
	ATOM	181	CE1	PHE	24	3.917	83.808	6.561	1.00	32.57
	ATOM	182	CE2	PHE	24	4.725	85.138	4.738	1.00	25.01
	ATOM	183	CZ	PHE	24	4.275	83.925	5.228	1.00	30.12
55	ATOM	184	C	PHE	24	6.063	88.675	9.319	1.00	31.73
	ATOM	185	O	PHE	24	5.393	88.948	10.310	1.00	33.31
	ATOM	186	N	VAL	25	6.955	89.506	8.805	1.00	32.92
	ATOM	187	CA	VAL	25	7.183	90.816	9.377	1.00	30.84
	ATOM	188	CB	VAL	25	8.410	91.487	8.741	1.00	30.29
60	ATOM	189	CG1	VAL	25	8.606	92.866	9.321	1.00	28.13
	ATOM	190	CG2	VAL	25	9.634	90.640	8.964	1.00	29.53
	ATOM	191	C	VAL	25	5.974	91.711	9.129	1.00	36.08
	ATOM	192	O	VAL	25	5.411	91.719	8.036	1.00	35.82
	ATOM	193	N	PRO	26	5.543	92.461	10.153	1.00	38.03
65	ATOM	194	CD	PRO	26	5.938	92.303	11.561	1.00	38.28
	ATOM	195	CA	PRO	26	4.396	93.373	10.040	1.00	38.08

	ATOM	196	CB	PRO	26	4.014	93.644	11.497	1.00	37.62
	ATOM	197	CG	PRO	26	4.634	92.517	12.260	1.00	36.31
	ATOM	198	C	PRO	26	4.929	94.638	9.369	1.00	40.85
	ATOM	199	O	PRO	26	5.720	95.368	9.965	1.00	42.27
5	ATOM	200	N	TRP	27	4.511	94.911	8.142	1.00	40.86
	ATOM	201	CA	TRP	27	5.035	96.084	7.454	1.00	37.05
	ATOM	202	CB	TRP	27	5.135	95.821	5.949	1.00	33.29
	ATOM	203	CG	TRP	27	6.033	94.688	5.605	1.00	31.06
	ATOM	204	CD2	TRP	27	7.431	94.587	5.882	1.00	27.79
10	ATOM	205	CE2	TRP	27	7.858	93.335	5.410	1.00	29.28
	ATOM	206	CE3	TRP	27	8.363	95.432	6.486	1.00	27.06
	ATOM	207	CD1	TRP	27	5.681	93.532	4.991	1.00	31.07
	ATOM	208	NE1	TRP	27	6.766	92.712	4.868	1.00	32.70
	ATOM	209	CZ2	TRP	27	9.181	92.901	5.521	1.00	25.99
15	ATOM	210	CZ3	TRP	27	9.680	95.001	6.596	1.00	26.93
	ATOM	211	CH2	TRP	27	10.075	93.746	6.115	1.00	28.08
	ATOM	212	C	TRP	27	4.274	97.371	7.676	1.00	38.50
	ATOM	213	O	TRP	27	3.103	97.370	8.040	1.00	39.07
	ATOM	214	N	LEU	28	4.982	98.473	7.450	1.00	40.19
20	ATOM	215	CA	LEU	28	4.440	99.817	7.572	1.00	39.45
	ATOM	216	CB	LEU	28	4.944	100.476	8.850	1.00	44.18
	ATOM	217	CG	LEU	28	4.174	101.714	9.307	1.00	46.83
	ATOM	218	CD1	LEU	28	2.767	101.289	9.736	1.00	45.15
	ATOM	219	CD2	LEU	28	4.917	102.389	10.464	1.00	49.15
25	ATOM	220	C	LEU	28	5.011	100.551	6.366	1.00	39.80
	ATOM	221	O	LEU	28	6.172	100.365	6.031	1.00	42.40
	ATOM	222	N	LEU	29	4.212	101.371	5.704	1.00	33.58
	ATOM	223	CA	LEU	29	4.719	102.079	4.546	1.00	29.21
	ATOM	224	CB	LEU	29	3.601	102.844	3.840	1.00	27.16
30	ATOM	225	CG	LEU	29	4.070	103.674	2.641	1.00	22.36
	ATOM	226	CD1	LEU	29	4.390	102.752	1.476	1.00	21.47
	ATOM	227	CD2	LEU	29	3.015	104.675	2.251	1.00	14.53
	ATOM	228	C	LEU	29	5.831	103.059	4.900	1.00	29.98
	ATOM	229	O	LEU	29	5.681	103.906	5.775	1.00	30.36
35	ATOM	230	N	SER	30	6.955	102.925	4.209	1.00	29.88
	ATOM	231	CA	SER	30	8.081	103.817	4.406	1.00	28.49
	ATOM	232	CB	SER	30	9.398	103.111	4.083	1.00	26.13
	ATOM	233	OG	SER	30	10.483	104.006	4.185	1.00	20.25
	ATOM	234	C	SER	30	7.834	104.957	3.427	1.00	31.61
40	ATOM	235	O	SER	30	7.827	106.125	3.803	1.00	35.14
	ATOM	236	N	PHE	31	7.622	104.604	2.164	1.00	32.12
	ATOM	237	CA	PHE	31	7.339	105.589	1.133	1.00	30.76
	ATOM	238	CB	PHE	31	8.567	106.467	0.864	1.00	27.55
	ATOM	239	CG	PHE	31	9.508	105.915	-0.170	1.00	31.23
45	ATOM	240	CD1	PHE	31	9.262	106.104	-1.525	1.00	32.00
	ATOM	241	CD2	PHE	31	10.637	105.193	0.210	1.00	27.89
	ATOM	242	CE1	PHE	31	10.121	105.585	-2.481	1.00	29.70
	ATOM	243	CE2	PHE	31	11.499	104.673	-0.742	1.00	26.15
	ATOM	244	CZ	PHE	31	11.241	104.868	-2.088	1.00	28.30
50	ATOM	245	C	PHE	31	6.900	104.890	-0.142	1.00	32.06
	ATOM	246	O	PHE	31	7.288	103.755	-0.415	1.00	32.26
	ATOM	247	N	LYS	32	6.072	105.574	-0.912	1.00	34.83
	ATOM	248	CA	LYS	32	5.588	105.039	-2.168	1.00	37.13
	ATOM	249	CB	LYS	32	4.102	104.700	-2.057	1.00	36.55
55	ATOM	250	CG	LYS	32	3.507	104.199	-3.343	1.00	40.62
	ATOM	251	CD	LYS	32	2.001	104.142	-3.285	1.00	44.56
	ATOM	252	CE	LYS	32	1.450	103.925	-4.696	1.00	46.29
	ATOM	253	NZ	LYS	32	-0.030	103.731	-4.725	1.00	50.58
	ATOM	254	C	LYS	32	5.814	106.117	-3.224	1.00	36.39
60	ATOM	255	O	LYS	32	5.511	107.289	-2.999	1.00	38.69
	ATOM	256	N	ARG	33	6.367	105.726	-4.367	1.00	37.90
	ATOM	257	CA	ARG	33	6.634	106.676	-5.437	1.00	35.74
	ATOM	258	CB	ARG	33	8.131	106.962	-5.491	1.00	36.79
	ATOM	259	CG	ARG	33	8.580	107.877	-6.608	1.00	31.09
65	ATOM	260	CD	ARG	33	10.028	108.291	-6.378	1.00	31.62
	ATOM	261	NE	ARG	33	10.501	109.296	-7.326	1.00	38.16

	ATOM	262	CZ	ARG	33	11.257	109.026	-8.384	1.00	39.15
	ATOM	263	NH1	ARG	33	11.633	107.780	-8.632	1.00	43.41
	ATOM	264	NH2	ARG	33	11.636	109.998	-9.196	1.00	40.65
	ATOM	265	C	ARG	33	6.150	106.115	-6.769	1.00	37.71
5	ATOM	266	O	ARG	33	6.610	105.061	-7.216	1.00	41.16
	ATOM	267	N	GLY	34	5.213	106.810	-7.406	1.00	36.59
	ATOM	268	CA	GLY	34	4.709	106.333	-8.676	1.00	36.72
	ATOM	269	C	GLY	34	3.477	105.466	-8.528	1.00	40.01
	ATOM	270	O	GLY	34	2.852	105.422	-7.469	1.00	39.85
10	ATOM	271	N	SER	35	3.144	104.751	-9.594	1.00	38.74
	ATOM	272	CA	SER	35	1.961	103.904	-9.610	1.00	40.23
	ATOM	273	CB	SER	35	1.029	104.392	-10.712	1.00	43.02
	ATOM	274	OG	SER	35	1.727	104.462	-11.954	1.00	48.05
	ATOM	275	C	SER	35	2.210	102.407	-9.811	1.00	40.54
15	ATOM	276	O	SER	35	1.329	101.597	-9.526	1.00	43.74
	ATOM	277	N	ALA	36	3.394	102.041	-10.296	1.00	35.56
	ATOM	278	CA	ALA	36	3.722	100.646	-10.568	1.00	29.62
	ATOM	279	CB	ALA	36	5.095	100.566	-11.201	1.00	25.47
	ATOM	280	C	ALA	36	3.645	99.685	-9.384	1.00	31.23
20	ATOM	281	O	ALA	36	3.445	98.492	-9.576	1.00	29.56
	ATOM	282	N	LEU	37	3.800	100.192	-8.166	1.00	34.09
	ATOM	283	CA	LEU	37	3.772	99.338	-6.983	1.00	32.32
	ATOM	284	CB	LEU	37	5.183	99.197	-6.415	1.00	26.47
	ATOM	285	CG	LEU	37	6.204	98.590	-7.374	1.00	25.72
25	ATOM	286	CD1	LEU	37	7.606	98.916	-6.924	1.00	24.61
	ATOM	287	CD2	LEU	37	5.992	97.097	-7.463	1.00	27.53
	ATOM	288	C	LEU	37	2.839	99.850	-5.899	1.00	35.16
	ATOM	289	O	LEU	37	2.733	101.049	-5.668	1.00	36.47
	ATOM	290	N	GLU	38	2.171	98.922	-5.232	1.00	38.30
30	ATOM	291	CA	GLU	38	1.244	99.258	-4.163	1.00	41.17
	ATOM	292	CB	GLU	38	-0.195	99.155	-4.656	1.00	46.15
	ATOM	293	CG	GLU	38	-0.763	100.409	-5.277	1.00	50.01
	ATOM	294	CD	GLU	38	-2.084	100.134	-5.985	1.00	51.18
	ATOM	295	OE1	GLU	38	-2.893	99.341	-5.438	1.00	48.40
35	ATOM	296	OE2	GLU	38	-2.307	100.711	-7.082	1.00	50.23
	ATOM	297	C	GLU	38	1.398	98.310	-2.989	1.00	42.69
	ATOM	298	O	GLU	38	1.880	97.195	-3.134	1.00	42.55
	ATOM	299	N	GLU	39	0.982	98.757	-1.815	1.00	44.75
	ATOM	300	CA	GLU	39	1.053	97.899	-0.644	1.00	46.50
40	ATOM	301	CB	GLU	39	1.356	98.721	0.605	1.00	47.52
	ATOM	302	CG	GLU	39	0.646	100.057	0.617	1.00	58.50
	ATOM	303	CD	GLU	39	0.839	100.807	1.921	1.00	62.35
	ATOM	304	OE1	GLU	39	0.467	102.007	1.984	1.00	64.58
	ATOM	305	OE2	GLU	39	1.353	100.191	2.885	1.00	63.04
45	ATOM	306	C	GLU	39	-0.331	97.285	-0.554	1.00	44.56
	ATOM	307	O	GLU	39	-1.327	97.967	-0.760	1.00	48.11
	ATOM	308	N	LYS	40	-0.399	95.995	-0.270	1.00	40.71
	ATOM	309	CA	LYS	40	-1.680	95.322	-0.178	1.00	38.08
	ATOM	310	CB	LYS	40	-2.114	94.825	-1.555	1.00	41.28
50	ATOM	311	CG	LYS	40	-3.361	93.949	-1.534	1.00	41.52
	ATOM	312	CD	LYS	40	-3.541	93.227	-2.862	1.00	42.73
	ATOM	313	CE	LYS	40	-4.682	92.226	-2.820	1.00	41.97
	ATOM	314	NZ	LYS	40	-4.754	91.446	-4.095	1.00	45.94
	ATOM	315	C	LYS	40	-1.614	94.149	0.778	1.00	39.66
55	ATOM	316	O	LYS	40	-0.926	93.164	0.524	1.00	36.18
	ATOM	317	N	GLU	41	-2.337	94.270	1.884	1.00	40.16
	ATOM	318	CA	GLU	41	-2.396	93.219	2.888	1.00	42.80
	ATOM	319	CB	GLU	41	-3.291	92.096	2.374	1.00	45.31
	ATOM	320	CG	GLU	41	-4.669	92.608	1.975	1.00	56.07
60	ATOM	321	CD	GLU	41	-5.438	91.644	1.076	1.00	59.42
	ATOM	322	OE1	GLU	41	-4.902	91.244	0.012	1.00	63.71
	ATOM	323	OE2	GLU	41	-6.584	91.290	1.432	1.00	63.71
	ATOM	324	C	GLU	41	-1.021	92.689	3.258	1.00	38.93
	ATOM	325	O	GLU	41	-0.768	91.489	3.196	1.00	39.03
65	ATOM	326	N	ASN	42	-0.135	93.605	3.625	1.00	34.69
	ATOM	327	CA	ASN	42	1.216	93.270	4.041	1.00	32.96

	ATOM	328	CB	ASN	42	1.175	92.265	5.186	1.00	32.65
	ATOM	329	CG	ASN	42	2.284	92.480	6.168	1.00	35.50
	ATOM	330	OD1	ASN	42	2.923	91.536	6.613	1.00	40.26
	ATOM	331	ND2	ASN	42	2.519	93.735	6.521	1.00	29.28
5	ATOM	332	C	ASN	42	2.121	92.728	2.945	1.00	31.44
	ATOM	333	O	ASN	42	3.151	92.124	3.229	1.00	22.78
	ATOM	334	N	LYS	43	1.740	92.952	1.698	1.00	33.26
	ATOM	335	CA	LYS	43	2.526	92.483	0.569	1.00	33.68
	ATOM	336	CB	LYS	43	1.848	91.279	-0.083	1.00	33.55
10	ATOM	337	CG	LYS	43	1.830	90.046	0.785	1.00	40.49
	ATOM	338	CD	LYS	43	1.065	88.929	0.121	1.00	45.39
	ATOM	339	CE	LYS	43	-0.401	89.289	-0.018	1.00	52.57
	ATOM	340	NZ	LYS	43	-1.184	88.156	-0.573	1.00	62.06
	ATOM	341	C	LYS	43	2.662	93.592	-0.449	1.00	32.05
15	ATOM	342	O	LYS	43	1.954	94.588	-0.380	1.00	37.32
	ATOM	343	N	ILE	44	3.582	93.433	-1.388	1.00	31.31
	ATOM	344	CA	ILE	44	3.747	94.446	-2.411	1.00	29.52
	ATOM	345	CB	ILE	44	5.224	94.716	-2.699	1.00	26.38
	ATOM	346	CG2	ILE	44	5.349	95.744	-3.805	1.00	26.09
20	ATOM	347	CG1	ILE	44	5.901	95.243	-1.434	1.00	28.61
	ATOM	348	CD1	ILE	44	7.354	95.570	-1.598	1.00	21.51
	ATOM	349	C	ILE	44	3.046	93.978	-3.674	1.00	31.60
	ATOM	350	O	ILE	44	3.340	92.908	-4.202	1.00	34.50
	ATOM	351	N	LEU	45	2.098	94.780	-4.140	1.00	31.44
25	ATOM	352	CA	LEU	45	1.338	94.463	-5.339	1.00	31.40
	ATOM	353	CB	LEU	45	-0.127	94.836	-5.130	1.00	33.60
	ATOM	354	CG	LEU	45	-1.041	94.672	-6.342	1.00	34.36
	ATOM	355	CD1	LEU	45	-1.178	93.206	-6.700	1.00	29.00
	ATOM	356	CD2	LEU	45	-2.394	95.276	-6.021	1.00	32.42
30	ATOM	357	C	LEU	45	1.876	95.183	-6.573	1.00	27.80
	ATOM	358	O	LEU	45	2.075	96.393	-6.565	1.00	27.94
	ATOM	359	N	VAL	46	2.108	94.420	-7.633	1.00	28.00
	ATOM	360	CA	VAL	46	2.608	94.966	-8.886	1.00	31.53
	ATOM	361	CB	VAL	46	3.408	93.895	-9.646	1.00	27.89
35	ATOM	362	CG1	VAL	46	3.975	94.466	-10.923	1.00	26.10
	ATOM	363	CG2	VAL	46	4.510	93.371	-8.767	1.00	23.39
	ATOM	364	C	VAL	46	1.427	95.434	-9.750	1.00	33.94
	ATOM	365	O	VAL	46	0.515	94.654	-10.036	1.00	39.68
	ATOM	366	N	LYS	47	1.444	96.698	-10.166	1.00	33.59
40	ATOM	367	CA	LYS	47	0.363	97.241	-10.983	1.00	37.05
	ATOM	368	CB	LYS	47	-0.129	98.569	-10.396	1.00	34.49
	ATOM	369	CG	LYS	47	-0.406	98.511	-8.904	1.00	37.53
	ATOM	370	CD	LYS	47	-1.426	97.433	-8.562	1.00	41.69
	ATOM	371	CE	LYS	47	-2.829	98.005	-8.481	1.00	42.95
45	ATOM	372	NZ	LYS	47	-3.128	98.880	-9.638	1.00	44.50
	ATOM	373	C	LYS	47	0.748	97.440	-12.450	1.00	37.34
	ATOM	374	O	LYS	47	-0.110	97.684	-13.292	1.00	43.04
	ATOM	375	N	GLU	48	2.036	97.338	-12.744	1.00	36.99
	ATOM	376	CA	GLU	48	2.542	97.483	-14.106	1.00	37.31
50	ATOM	377	CB	GLU	48	3.221	98.829	-14.297	1.00	36.53
	ATOM	378	CG	GLU	48	2.308	100.019	-14.273	1.00	46.74
	ATOM	379	CD	GLU	48	3.084	101.315	-14.423	1.00	52.39
	ATOM	380	OE1	GLU	48	4.042	101.340	-15.244	1.00	51.33
	ATOM	381	OE2	GLU	48	2.736	102.304	-13.727	1.00	54.55
55	ATOM	382	C	GLU	48	3.583	96.410	-14.306	1.00	36.94
	ATOM	383	O	GLU	48	4.493	96.293	-13.495	1.00	40.21
	ATOM	384	N	THR	49	3.485	95.631	-15.375	1.00	36.34
	ATOM	385	CA	THR	49	4.479	94.585	-15.556	1.00	35.48
	ATOM	386	CB	THR	49	3.981	93.467	-16.514	1.00	31.16
60	ATOM	387	OG1	THR	49	4.484	93.699	-17.828	1.00	32.48
	ATOM	388	CG2	THR	49	2.473	93.434	-16.561	1.00	30.36
	ATOM	389	C	THR	49	5.796	95.178	-16.066	1.00	34.59
	ATOM	390	O	THR	49	5.811	96.203	-16.744	1.00	35.13
	ATOM	391	N	GLY	50	6.901	94.528	-15.709	1.00	32.49
65	ATOM	392	CA	GLY	50	8.205	94.993	-16.128	1.00	28.17
	ATOM	393	C	GLY	50	9.297	94.360	-15.297	1.00	29.99

	ATOM	394	O	GLY	50	9.076	93.345	-14.656	1.00	29.76
	ATOM	395	N	TYR	51	10.483	94.958	-15.318	1.00	31.68
	ATOM	396	CA	TYR	51	11.616	94.461	-14.545	1.00	32.88
	ATOM	397	CB	TYR	51	12.899	94.495	-15.383	1.00	35.97
5	ATOM	398	CG	TYR	51	12.907	93.471	-16.480	1.00	43.03
	ATOM	399	CD1	TYR	51	12.244	93.708	-17.688	1.00	44.15
	ATOM	400	CE1	TYR	51	12.169	92.723	-18.675	1.00	47.81
	ATOM	401	CD2	TYR	51	13.504	92.229	-16.283	1.00	47.08
	ATOM	402	CE2	TYR	51	13.435	91.233	-17.257	1.00	51.79
10	ATOM	403	CZ	TYR	51	12.763	91.483	-18.451	1.00	51.15
	ATOM	404	OH	TYR	51	12.652	90.478	-19.393	1.00	52.20
	ATOM	405	C	TYR	51	11.808	95.282	-13.277	1.00	31.97
	ATOM	406	O	TYR	51	11.910	96.506	-13.324	1.00	30.01
	ATOM	407	N	PHE	52	11.855	94.600	-12.139	1.00	31.50
15	ATOM	408	CA	PHE	52	12.020	95.284	-10.867	1.00	30.41
	ATOM	409	CB	PHE	52	10.786	95.080	-9.985	1.00	31.30
	ATOM	410	CG	PHE	52	9.509	95.575	-10.586	1.00	29.70
	ATOM	411	CD1	PHE	52	8.899	94.888	-11.622	1.00	26.20
	ATOM	412	CD2	PHE	52	8.908	96.729	-10.104	1.00	28.38
20	ATOM	413	CE1	PHE	52	7.716	95.342	-12.167	1.00	27.17
	ATOM	414	CE2	PHE	52	7.722	97.191	-10.645	1.00	27.76
	ATOM	415	CZ	PHE	52	7.125	96.498	-11.676	1.00	27.27
	ATOM	416	C	PHE	52	13.231	94.823	-10.083	1.00	27.65
	ATOM	417	O	PHE	52	13.622	93.665	-10.143	1.00	24.85
25	ATOM	418	N	PHE	53	13.818	95.756	-9.350	1.00	27.18
	ATOM	419	CA	PHE	53	14.946	95.467	-8.478	1.00	27.05
	ATOM	420	CB	PHE	53	15.906	96.648	-8.418	1.00	28.76
	ATOM	421	CG	PHE	53	16.984	96.494	-7.394	1.00	28.00
	ATOM	422	CD1	PHE	53	18.008	95.576	-7.572	1.00	26.49
30	ATOM	423	CD2	PHE	53	16.967	97.256	-6.235	1.00	28.04
	ATOM	424	CE1	PHE	53	19.001	95.422	-6.611	1.00	26.54
	ATOM	425	CE2	PHE	53	17.959	97.103	-5.270	1.00	25.78
	ATOM	426	CZ	PHE	53	18.974	96.185	-5.462	1.00	24.39
	ATOM	427	C	PHE	53	14.265	95.293	-7.122	1.00	27.56
35	ATOM	428	O	PHE	53	13.599	96.206	-6.645	1.00	25.39
	ATOM	429	N	ILE	54	14.421	94.119	-6.519	1.00	28.87
	ATOM	430	CA	ILE	54	13.784	93.825	-5.239	1.00	28.95
	ATOM	431	CB	ILE	54	12.886	92.579	-5.379	1.00	29.60
	ATOM	432	CG2	ILE	54	12.010	92.428	-4.146	1.00	27.42
40	ATOM	433	CG1	ILE	54	12.012	92.719	-6.627	1.00	26.41
	ATOM	434	CD1	ILE	54	11.339	91.453	-7.063	1.00	28.70
	ATOM	435	C	ILE	54	14.820	93.596	-4.138	1.00	28.57
	ATOM	436	O	ILE	54	15.824	92.923	-4.346	1.00	29.17
	ATOM	437	N	TYR	55	14.576	94.153	-2.959	1.00	26.97
45	ATOM	438	CA	TYR	55	15.529	94.008	-1.866	1.00	24.22
	ATOM	439	CB	TYR	55	16.447	95.231	-1.823	1.00	21.82
	ATOM	440	CG	TYR	55	15.724	96.543	-1.642	1.00	27.68
	ATOM	441	CD1	TYR	55	15.481	97.059	-0.373	1.00	27.01
	ATOM	442	CE1	TYR	55	14.799	98.252	-0.208	1.00	27.44
50	ATOM	443	CD2	TYR	55	15.263	97.260	-2.742	1.00	27.16
	ATOM	444	CE2	TYR	55	14.576	98.453	-2.584	1.00	25.39
	ATOM	445	CZ	TYR	55	14.348	98.942	-1.319	1.00	29.02
	ATOM	446	OH	TYR	55	13.658	100.117	-1.165	1.00	31.82
	ATOM	447	C	TYR	55	14.868	93.807	-0.520	1.00	25.16
55	ATOM	448	O	TYR	55	13.697	94.069	-0.358	1.00	28.47
	ATOM	449	N	GLY	56	15.627	93.327	0.451	1.00	27.44
	ATOM	450	CA	GLY	56	15.069	93.105	1.767	1.00	22.07
	ATOM	451	C	GLY	56	16.135	92.805	2.798	1.00	25.83
	ATOM	452	O	GLY	56	17.152	92.194	2.486	1.00	26.00
60	ATOM	453	N	GLN	57	15.909	93.260	4.024	1.00	22.56
	ATOM	454	CA	GLN	57	16.832	93.024	5.122	1.00	20.91
	ATOM	455	CB	GLN	57	17.770	94.211	5.341	1.00	17.87
	ATOM	456	CG	GLN	57	18.655	94.040	6.570	1.00	16.86
	ATOM	457	CD	GLN	57	19.653	95.162	6.765	1.00	22.06
65	ATOM	458	OE1	GLN	57	20.546	95.359	5.949	1.00	27.26
	ATOM	459	NE2	GLN	57	19.507	95.902	7.858	1.00	20.20

	ATOM	460	C	GLN	57	16.038	92.781	6.388	1.00	24.23
	ATOM	461	O	GLN	57	14.984	93.383	6.589	1.00	25.16
	ATOM	462	N	VAL	58	16.555	91.894	7.233	1.00	24.62
	ATOM	463	CA	VAL	58	15.929	91.544	8.503	1.00	26.68
5	ATOM	464	CB	VAL	58	15.172	90.197	8.406	1.00	25.55
	ATOM	465	CG1	VAL	58	14.715	89.757	9.776	1.00	15.69
	ATOM	466	CG2	VAL	58	13.982	90.324	7.479	1.00	25.77
	ATOM	467	C	VAL	58	17.030	91.382	9.530	1.00	31.06
	ATOM	468	O	VAL	58	18.081	90.845	9.208	1.00	27.74
10	ATOM	469	N	LEU	59	16.798	91.848	10.757	1.00	33.82
	ATOM	470	CA	LEU	59	17.779	91.699	11.838	1.00	29.85
	ATOM	471	CB	LEU	59	17.839	92.948	12.713	1.00	29.62
	ATOM	472	CG	LEU	59	19.129	93.190	13.510	1.00	28.11
	ATOM	473	CD1	LEU	59	18.871	94.248	14.563	1.00	26.67
15	ATOM	474	CD2	LEU	59	19.617	91.931	14.163	1.00	22.45
	ATOM	475	C	LEU	59	17.347	90.517	12.700	1.00	30.98
	ATOM	476	O	LEU	59	16.302	90.557	13.346	1.00	32.52
	ATOM	477	N	TYR	60	18.154	89.460	12.705	1.00	33.01
	ATOM	478	CA	TYR	60	17.838	88.276	13.490	1.00	34.51
20	ATOM	479	CB	TYR	60	18.306	87.022	12.757	1.00	40.10
	ATOM	480	CG	TYR	60	17.620	86.840	11.439	1.00	45.37
	ATOM	481	CD1	TYR	60	18.289	87.088	10.239	1.00	47.13
	ATOM	482	CE1	TYR	60	17.624	87.001	9.024	1.00	50.28
	ATOM	483	CD2	TYR	60	16.276	86.494	11.388	1.00	44.35
25	ATOM	484	CE2	TYR	60	15.602	86.403	10.187	1.00	48.81
	ATOM	485	CZ	TYR	60	16.276	86.662	9.011	1.00	51.91
	ATOM	486	OH	TYR	60	15.584	86.612	7.826	1.00	53.77
	ATOM	487	C	TYR	60	18.440	88.302	14.884	1.00	35.49
	ATOM	488	O	TYR	60	19.649	88.424	15.054	1.00	36.13
30	ATOM	489	N	THR	61	17.576	88.189	15.883	1.00	36.19
	ATOM	490	CA	THR	61	18.002	88.178	17.269	1.00	35.36
	ATOM	491	CB	THR	61	17.396	89.356	18.034	1.00	34.22
	ATOM	492	OG1	THR	61	15.977	89.363	17.851	1.00	34.22
	ATOM	493	CG2	THR	61	17.976	90.667	17.536	1.00	25.45
35	ATOM	494	C	THR	61	17.519	86.864	17.866	1.00	37.82
	ATOM	495	O	THR	61	17.178	86.777	19.042	1.00	43.70
	ATOM	496	N	ASP	62	17.494	85.844	17.019	1.00	37.64
	ATOM	497	CA	ASP	62	17.060	84.505	17.374	1.00	38.08
	ATOM	498	CB	ASP	62	16.074	84.035	16.308	1.00	38.42
40	ATOM	499	CG	ASP	62	15.409	82.734	16.655	1.00	41.25
	ATOM	500	OD1	ASP	62	14.194	82.604	16.372	1.00	38.28
	ATOM	501	OD2	ASP	62	16.103	81.848	17.194	1.00	41.37
	ATOM	502	C	ASP	62	18.301	83.618	17.398	1.00	40.90
	ATOM	503	O	ASP	62	19.177	83.780	16.559	1.00	46.73
45	ATOM	504	N	LYS	63	18.397	82.686	18.342	1.00	42.62
	ATOM	505	CA	LYS	63	19.588	81.838	18.401	1.00	40.92
	ATOM	506	CB	LYS	63	20.056	81.678	19.849	1.00	42.79
	ATOM	507	CG	LYS	63	19.034	81.071	20.786	1.00	45.91
	ATOM	508	CD	LYS	63	19.522	81.159	22.232	1.00	46.22
50	ATOM	509	CE	LYS	63	19.707	82.626	22.666	1.00	49.97
	ATOM	510	NZ	LYS	63	20.394	82.789	23.993	1.00	49.98
	ATOM	511	C	LYS	63	19.448	80.464	17.754	1.00	42.59
	ATOM	512	O	LYS	63	20.233	79.566	18.031	1.00	42.46
	ATOM	513	N	THR	64	18.475	80.323	16.863	1.00	44.06
55	ATOM	514	CA	THR	64	18.221	79.064	16.177	1.00	44.94
	ATOM	515	CB	THR	64	16.859	79.123	15.440	1.00	43.79
	ATOM	516	OG1	THR	64	15.833	79.475	16.374	1.00	46.20
	ATOM	517	CG2	THR	64	16.507	77.774	14.825	1.00	42.80
	ATOM	518	C	THR	64	19.301	78.595	15.189	1.00	45.82
60	ATOM	519	O	THR	64	18.986	78.150	14.100	1.00	53.10
	ATOM	520	N	TYR	65	20.567	78.680	15.566	1.00	45.11
	ATOM	521	CA	TYR	65	21.687	78.227	14.728	1.00	45.72
	ATOM	522	CB	TYR	65	21.774	76.689	14.751	1.00	40.46
	ATOM	523	CG	TYR	65	21.068	75.970	13.623	1.00	41.90
65	ATOM	524	CD1	TYR	65	21.714	75.723	12.409	1.00	42.47
	ATOM	525	CE1	TYR	65	21.058	75.064	11.354	1.00	41.83

	ATOM	526	CD2	TYR	65	19.745	75.540	13.761	1.00	41.65
	ATOM	527	CE2	TYR	65	19.076	74.882	12.714	1.00	40.28
	ATOM	528	CZ	TYR	65	19.742	74.650	11.517	1.00	41.90
	ATOM	529	OH	TYR	65	19.094	74.008	10.491	1.00	43.84
5	ATOM	530	C	TYR	65	21.746	78.724	13.274	1.00	45.31
	ATOM	531	O	TYR	65	22.836	78.878	12.708	1.00	49.74
	ATOM	532	N	ALA	66	20.593	78.965	12.667	1.00	42.32
	ATOM	533	CA	ALA	66	20.537	79.439	11.290	1.00	39.57
	ATOM	534	CB	ALA	66	20.705	78.281	10.331	1.00	35.36
10	ATOM	535	C	ALA	66	19.213	80.136	11.040	1.00	39.85
	ATOM	536	O	ALA	66	18.144	79.547	11.214	1.00	37.60
	ATOM	537	N	MET	67	19.287	81.402	10.649	1.00	41.88
	ATOM	538	CA	MET	67	18.091	82.183	10.369	1.00	41.48
	ATOM	539	CB	MET	67	17.941	83.333	11.372	1.00	39.98
15	ATOM	540	CG	MET	67	17.620	82.903	12.798	1.00	37.31
	ATOM	541	SD	MET	67	16.093	81.963	12.925	1.00	35.46
	ATOM	542	CE	MET	67	14.860	83.267	12.917	1.00	34.06
	ATOM	543	C	MET	67	18.175	82.737	8.958	1.00	39.50
	ATOM	544	O	MET	67	19.239	82.719	8.334	1.00	38.44
20	ATOM	545	N	GLY	68	17.048	83.226	8.458	1.00	39.84
	ATOM	546	CA	GLY	68	17.018	83.782	7.120	1.00	37.61
	ATOM	547	C	GLY	68	15.611	84.063	6.653	1.00	33.76
	ATOM	548	O	GLY	68	14.649	83.705	7.323	1.00	35.58
	ATOM	549	N	HIS	69	15.486	84.721	5.509	1.00	35.07
25	ATOM	550	CA	HIS	69	14.177	85.035	4.960	1.00	33.70
	ATOM	551	CB	HIS	69	13.801	86.502	5.240	1.00	37.02
	ATOM	552	CG	HIS	69	14.823	87.501	4.794	1.00	34.65
	ATOM	553	CD2	HIS	69	14.834	88.354	3.745	1.00	35.63
	ATOM	554	ND1	HIS	69	16.003	87.718	5.472	1.00	37.68
30	ATOM	555	CE1	HIS	69	16.696	88.661	4.862	1.00	32.64
	ATOM	556	NE2	HIS	69	16.008	89.065	3.811	1.00	35.90
	ATOM	557	C	HIS	69	14.115	84.755	3.466	1.00	33.98
	ATOM	558	O	HIS	69	15.127	84.500	2.833	1.00	30.94
	ATOM	559	N	LEU	70	12.912	84.798	2.909	1.00	35.35
35	ATOM	560	CA	LEU	70	12.712	84.550	1.490	1.00	30.76
	ATOM	561	CB	LEU	70	11.897	83.278	1.284	1.00	29.59
	ATOM	562	CG	LEU	70	12.172	82.069	2.173	1.00	30.71
	ATOM	563	CD1	LEU	70	11.086	81.042	1.962	1.00	25.60
	ATOM	564	CD2	LEU	70	13.534	81.496	1.862	1.00	27.39
40	ATOM	565	C	LEU	70	11.927	85.697	0.861	1.00	33.19
	ATOM	566	O	LEU	70	10.974	86.203	1.453	1.00	37.07
	ATOM	567	N	ILE	71	12.337	86.130	-0.326	1.00	30.93
	ATOM	568	CA	ILE	71	11.587	87.161	-1.025	1.00	27.94
	ATOM	569	CB	ILE	71	12.498	88.167	-1.733	1.00	28.22
45	ATOM	570	CG2	ILE	71	11.696	88.998	-2.712	1.00	24.54
	ATOM	571	CG1	ILE	71	13.144	89.082	-0.700	1.00	26.34
	ATOM	572	CD1	ILE	71	14.208	89.978	-1.270	1.00	37.50
	ATOM	573	C	ILE	71	10.840	86.317	-2.045	1.00	30.32
	ATOM	574	O	ILE	71	11.459	85.696	-2.902	1.00	23.07
50	ATOM	575	N	GLN	72	9.516	86.267	-1.935	1.00	29.74
	ATOM	576	CA	GLN	72	8.738	85.438	-2.844	1.00	27.85
	ATOM	577	CB	GLN	72	7.931	84.433	-2.036	1.00	26.70
	ATOM	578	CG	GLN	72	8.771	83.693	-1.035	1.00	28.28
	ATOM	579	CD	GLN	72	8.022	82.568	-0.380	1.00	33.11
55	ATOM	580	OE1	GLN	72	6.942	82.771	0.168	1.00	41.22
	ATOM	581	NE2	GLN	72	8.590	81.371	-0.424	1.00	30.18
	ATOM	582	C	GLN	72	7.817	86.179	-3.789	1.00	29.85
	ATOM	583	O	GLN	72	7.423	87.319	-3.536	1.00	28.20
	ATOM	584	N	ARG	73	7.468	85.502	-4.877	1.00	30.90
60	ATOM	585	CA	ARG	73	6.585	86.058	-5.889	1.00	30.79
	ATOM	586	CB	ARG	73	7.348	86.207	-7.199	1.00	29.18
	ATOM	587	CG	ARG	73	6.507	86.690	-8.341	1.00	28.66
	ATOM	588	CD	ARG	73	7.128	86.286	-9.646	1.00	29.25
	ATOM	589	NE	ARG	73	6.365	86.783	-10.780	1.00	35.78
65	ATOM	590	CZ	ARG	73	6.466	86.306	-12.011	1.00	35.27
	ATOM	591	NH1	ARG	73	7.298	85.307	-12.270	1.00	34.95

	ATOM	592	NH2	ARG	73	5.738	86.831	-12.980	1.00	33.33
	ATOM	593	C	ARG	73	5.361	85.175	-6.123	1.00	32.42
	ATOM	594	O	ARG	73	5.494	83.977	-6.348	1.00	34.19
	ATOM	595	N	LYS	74	4.174	85.770	-6.049	1.00	35.90
5	ATOM	596	CA	LYS	74	2.931	85.044	-6.307	1.00	38.57
	ATOM	597	CB	LYS	74	1.833	85.448	-5.320	1.00	43.52
	ATOM	598	CG	LYS	74	2.016	84.918	-3.908	1.00	52.89
	ATOM	599	CD	LYS	74	0.900	85.401	-2.952	1.00	58.14
	ATOM	600	CE	LYS	74	1.122	84.863	-1.537	1.00	56.75
10	ATOM	601	NZ	LYS	74	0.084	85.348	-0.579	1.00	63.98
	ATOM	602	C	LYS	74	2.478	85.417	-7.713	1.00	40.11
	ATOM	603	O	LYS	74	1.994	86.533	-7.939	1.00	37.55
	ATOM	604	N	LYS	75	2.634	84.490	-8.652	1.00	34.48
	ATOM	605	CA	LYS	75	2.250	84.731	-10.038	1.00	34.24
15	ATOM	606	CB	LYS	75	2.733	83.578	-10.921	1.00	36.16
	ATOM	607	CG	LYS	75	4.231	83.281	-10.899	1.00	35.99
	ATOM	608	CD	LYS	75	4.531	82.093	-11.823	1.00	40.91
	ATOM	609	CE	LYS	75	6.016	81.761	-11.866	1.00	46.16
	ATOM	610	NZ	LYS	75	6.366	80.568	-12.713	1.00	46.03
20	ATOM	611	C	LYS	75	0.731	84.861	-10.187	1.00	30.92
	ATOM	612	O	LYS	75	-0.011	84.127	-9.561	1.00	27.47
	ATOM	613	N	VAL	76	0.280	85.793	-11.023	1.00	32.82
	ATOM	614	CA	VAL	76	-1.152	85.984	-11.274	1.00	31.51
	ATOM	615	CB	VAL	76	-1.471	87.326	-11.957	1.00	29.19
25	ATOM	616	CG1	VAL	76	-2.790	87.846	-11.462	1.00	27.09
	ATOM	617	CG2	VAL	76	-0.378	88.308	-11.729	1.00	34.94
	ATOM	618	C	VAL	76	-1.586	84.921	-12.262	1.00	33.40
	ATOM	619	O	VAL	76	-2.666	84.351	-12.147	1.00	32.92
	ATOM	620	N	HIS	77	-0.730	84.683	-13.250	1.00	34.88
30	ATOM	621	CA	HIS	77	-0.984	83.703	-14.290	1.00	34.59
	ATOM	622	CB	HIS	77	-0.697	84.312	-15.667	1.00	35.92
	ATOM	623	CG	HIS	77	-1.498	85.542	-15.964	1.00	34.17
	ATOM	624	CD2	HIS	77	-1.256	86.584	-16.791	1.00	31.44
	ATOM	625	ND1	HIS	77	-2.732	85.785	-15.396	1.00	35.05
35	ATOM	626	CE1	HIS	77	-3.215	86.923	-15.857	1.00	29.84
	ATOM	627	NE2	HIS	77	-2.340	87.426	-16.707	1.00	38.35
	ATOM	628	C	HIS	77	-0.107	82.485	-14.052	1.00	35.63
	ATOM	629	O	HIS	77	1.064	82.621	-13.698	1.00	40.24
	ATOM	630	N	VAL	78	-0.664	81.295	-14.271	1.00	37.37
40	ATOM	631	CA	VAL	78	0.085	80.079	-14.014	1.00	41.32
	ATOM	632	CB	VAL	78	-0.594	79.279	-12.885	1.00	43.10
	ATOM	633	CG1	VAL	78	0.253	78.088	-12.512	1.00	45.01
	ATOM	634	CG2	VAL	78	-0.767	80.162	-11.662	1.00	44.98
	ATOM	635	C	VAL	78	0.450	79.134	-15.166	1.00	43.02
45	ATOM	636	O	VAL	78	1.632	78.994	-15.475	1.00	49.79
	ATOM	637	N	PHE	79	-0.509	78.472	-15.800	1.00	41.33
	ATOM	638	CA	PHE	79	-0.162	77.541	-16.902	1.00	45.00
	ATOM	639	CB	PHE	79	0.927	78.099	-17.840	1.00	40.51
	ATOM	640	CG	PHE	79	0.613	79.433	-18.428	1.00	39.52
50	ATOM	641	CD1	PHE	79	1.251	80.575	-17.957	1.00	36.80
	ATOM	642	CD2	PHE	79	-0.313	79.551	-19.455	1.00	37.89
	ATOM	643	CE1	PHE	79	0.972	81.818	-18.501	1.00	37.52
	ATOM	644	CE2	PHE	79	-0.598	80.791	-20.005	1.00	35.59
	ATOM	645	CZ	PHE	79	0.045	81.927	-19.528	1.00	37.58
55	ATOM	646	C	PHE	79	0.363	76.180	-16.443	1.00	43.50
	ATOM	647	O	PHE	79	1.348	76.092	-15.708	1.00	37.85
	ATOM	648	N	GLY	80	-0.286	75.123	-16.919	1.00	46.65
	ATOM	649	CA	GLY	80	0.124	73.770	-16.602	1.00	48.38
	ATOM	650	C	GLY	80	0.436	73.453	-15.155	1.00	47.89
60	ATOM	651	O	GLY	80	-0.339	73.781	-14.252	1.00	51.15
	ATOM	652	N	ASP	81	1.581	72.811	-14.938	1.00	45.07
	ATOM	653	CA	ASP	81	1.990	72.415	-13.603	1.00	43.99
	ATOM	654	CB	ASP	81	2.618	71.015	-13.640	1.00	45.66
	ATOM	655	CG	ASP	81	3.964	70.987	-14.348	1.00	47.49
65	ATOM	656	OD1	ASP	81	4.312	71.967	-15.044	1.00	51.90
	ATOM	657	OD2	ASP	81	4.677	69.973	-14.217	1.00	48.08

	ATOM	658	C	ASP	81	2.936	73.390	-12.925	1.00	42.29
	ATOM	659	O	ASP	81	3.679	73.008	-12.022	1.00	39.94
	ATOM	660	N	GLU	82	2.912	74.648	-13.357	1.00	39.91
	ATOM	661	CA	GLU	82	3.758	75.669	-12.743	1.00	36.58
5	ATOM	662	CB	GLU	82	3.608	77.023	-13.438	1.00	34.91
	ATOM	663	CG	GLU	82	4.341	77.223	-14.725	1.00	37.82
	ATOM	664	CD	GLU	82	4.607	78.697	-14.986	1.00	40.75
	ATOM	665	OE1	GLU	82	3.694	79.519	-14.771	1.00	38.15
	ATOM	666	OE2	GLU	82	5.733	79.041	-15.402	1.00	47.99
10	ATOM	667	C	GLU	82	3.322	75.881	-11.300	1.00	36.58
	ATOM	668	O	GLU	82	2.148	75.731	-10.972	1.00	38.40
	ATOM	669	N	LEU	83	4.270	76.228	-10.442	1.00	33.83
	ATOM	670	CA	LEU	83	3.962	76.537	-9.055	1.00	32.70
	ATOM	671	CB	LEU	83	5.169	76.269	-8.157	1.00	30.78
15	ATOM	672	CG	LEU	83	5.274	74.900	-7.488	1.00	29.26
	ATOM	673	CD1	LEU	83	4.916	73.799	-8.455	1.00	29.60
	ATOM	674	CD2	LEU	83	6.676	74.718	-6.965	1.00	28.65
	ATOM	675	C	LEU	83	3.697	78.026	-9.129	1.00	33.88
	ATOM	676	O	LEU	83	4.468	78.753	-9.745	1.00	38.46
20	ATOM	677	N	SER	84	2.611	78.487	-8.525	1.00	34.94
	ATOM	678	CA	SER	84	2.295	79.906	-8.585	1.00	39.77
	ATOM	679	CB	SER	84	0.799	80.128	-8.371	1.00	38.92
	ATOM	680	OG	SER	84	0.396	79.608	-7.128	1.00	43.11
	ATOM	681	C	SER	84	3.093	80.745	-7.591	1.00	39.89
25	ATOM	682	O	SER	84	3.095	81.973	-7.656	1.00	42.52
	ATOM	683	N	LEU	85	3.776	80.075	-6.674	1.00	40.92
	ATOM	684	CA	LEU	85	4.584	80.759	-5.683	1.00	34.66
	ATOM	685	CB	LEU	85	4.215	80.273	-4.282	1.00	35.12
	ATOM	686	CG	LEU	85	4.705	81.002	-3.023	1.00	36.92
30	ATOM	687	CD1	LEU	85	6.198	80.845	-2.873	1.00	38.24
	ATOM	688	CD2	LEU	85	4.316	82.471	-3.095	1.00	36.08
	ATOM	689	C	LEU	85	6.035	80.442	-5.994	1.00	35.30
	ATOM	690	O	LEU	85	6.464	79.294	-5.926	1.00	32.85
	ATOM	691	N	VAL	86	6.786	81.468	-6.364	1.00	33.26
35	ATOM	692	CA	VAL	86	8.194	81.303	-6.686	1.00	34.57
	ATOM	693	CB	VAL	86	8.535	81.859	-8.099	1.00	35.39
	ATOM	694	CG1	VAL	86	10.033	81.827	-8.329	1.00	31.44
	ATOM	695	CG2	VAL	86	7.831	81.042	-9.164	1.00	38.08
	ATOM	696	C	VAL	86	9.022	82.065	-5.680	1.00	32.85
40	ATOM	697	O	VAL	86	8.724	83.215	-5.368	1.00	35.10
	ATOM	698	N	THR	87	10.057	81.435	-5.153	1.00	33.41
	ATOM	699	CA	THR	87	10.904	82.148	-4.227	1.00	33.56
	ATOM	700	CB	THR	87	11.200	81.304	-2.949	1.00	32.70
	ATOM	701	OG1	THR	87	12.569	81.466	-2.556	1.00	34.29
45	ATOM	702	CG2	THR	87	10.872	79.852	-3.176	1.00	34.95
	ATOM	703	C	THR	87	12.188	82.603	-4.932	1.00	30.57
	ATOM	704	O	THR	87	12.982	81.792	-5.403	1.00	28.08
	ATOM	705	N	LEU	88	12.328	83.919	-5.051	1.00	28.42
	ATOM	706	CA	LEU	88	13.495	84.540	-5.648	1.00	31.31
50	ATOM	707	CB	LEU	88	13.134	85.879	-6.306	1.00	28.21
	ATOM	708	CG	LEU	88	12.048	86.199	-7.338	1.00	29.12
	ATOM	709	CD1	LEU	88	11.406	84.960	-7.857	1.00	31.12
	ATOM	710	CD2	LEU	88	11.024	87.112	-6.708	1.00	28.72
	ATOM	711	C	LEU	88	14.443	84.855	-4.478	1.00	41.15
55	ATOM	712	O	LEU	88	14.003	85.130	-3.359	1.00	52.30
	ATOM	713	N	PHE	89	15.742	84.799	-4.702	1.00	39.23
	ATOM	714	CA	PHE	89	16.667	85.182	-3.633	1.00	42.27
	ATOM	715	CB	PHE	89	16.439	86.667	-3.340	1.00	35.63
	ATOM	716	CG	PHE	89	16.238	87.477	-4.594	1.00	35.06
60	ATOM	717	CD1	PHE	89	15.237	88.439	-4.669	1.00	31.56
	ATOM	718	CD2	PHE	89	16.980	87.180	-5.751	1.00	30.96
	ATOM	719	CE1	PHE	89	14.963	89.085	-5.879	1.00	33.76
	ATOM	720	CE2	PHE	89	16.715	87.815	-6.960	1.00	27.38
	ATOM	721	CZ	PHE	89	15.704	88.765	-7.027	1.00	30.29
65	ATOM	722	C	PHE	89	16.715	84.341	-2.338	1.00	39.99
	ATOM	723	O	PHE	89	17.073	83.160	-2.386	1.00	45.34

	ATOM	724	N	ARG	90	16.393	84.909	-1.185	1.00	38.92
	ATOM	725	CA	ARG	90	16.500	84.124	0.059	1.00	41.66
	ATOM	726	CB	ARG	90	15.971	82.689	-0.145	1.00	39.24
	ATOM	727	CG	ARG	90	16.749	81.595	0.614	1.00	29.41
5	ATOM	728	CD	ARG	90	16.155	80.191	0.391	1.00	31.13
	ATOM	729	NE	ARG	90	17.040	79.129	0.876	1.00	31.75
	ATOM	730	CZ	ARG	90	17.653	78.230	0.102	1.00	28.86
	ATOM	731	NH1	ARG	90	17.487	78.232	-1.210	1.00	25.48
	ATOM	732	NH2	ARG	90	18.466	77.338	0.643	1.00	29.84
10	ATOM	733	C	ARG	90	17.930	84.020	0.624	1.00	39.23
	ATOM	734	O	ARG	90	18.865	83.666	-0.091	1.00	31.33
	ATOM	735	N	CYS	91	18.090	84.331	1.913	1.00	40.12
	ATOM	736	CA	CYS	91	19.392	84.216	2.551	1.00	37.09
	ATOM	737	C	CYS	91	19.418	83.585	3.914	1.00	37.21
15	ATOM	738	O	CYS	91	18.386	83.410	4.551	1.00	37.46
	ATOM	739	CB	CYS	91	20.113	85.542	2.629	1.00	40.13
	ATOM	740	SG	CYS	91	19.451	86.973	3.553	1.00	41.64
	ATOM	741	N	ILE	92	20.627	83.252	4.353	1.00	34.02
	ATOM	742	CA	ILE	92	20.831	82.592	5.634	1.00	33.22
20	ATOM	743	CB	ILE	92	21.122	81.093	5.417	1.00	32.88
	ATOM	744	CG2	ILE	92	21.144	80.367	6.745	1.00	32.61
	ATOM	745	CG1	ILE	92	20.049	80.485	4.512	1.00	34.64
	ATOM	746	CD1	ILE	92	20.334	79.060	4.067	1.00	32.30
	ATOM	747	C	ILE	92	21.982	83.189	6.433	1.00	30.78
25	ATOM	748	O	ILE	92	22.901	83.751	5.874	1.00	33.43
	ATOM	749	N	GLN	93	21.906	83.084	7.755	1.00	29.35
	ATOM	750	CA	GLN	93	22.952	83.571	8.639	1.00	28.69
	ATOM	751	CB	GLN	93	22.624	84.953	9.206	1.00	28.17
	ATOM	752	CG	GLN	93	22.980	86.166	8.343	1.00	25.87
30	ATOM	753	CD	GLN	93	24.456	86.287	8.034	1.00	24.22
	ATOM	754	OE1	GLN	93	24.934	85.740	7.052	1.00	28.90
	ATOM	755	NE2	GLN	93	25.186	87.005	8.876	1.00	25.29
	ATOM	756	C	GLN	93	23.042	82.604	9.799	1.00	31.44
	ATOM	757	O	GLN	93	22.039	82.314	10.433	1.00	32.07
35	ATOM	758	N	ASN	94	24.228	82.075	10.062	1.00	35.77
	ATOM	759	CA	ASN	94	24.387	81.186	11.198	1.00	33.56
	ATOM	760	CB	ASN	94	25.755	80.502	11.173	1.00	34.70
	ATOM	761	CG	ASN	94	25.792	79.304	10.259	1.00	33.01
	ATOM	762	OD1	ASN	94	24.954	78.423	10.352	1.00	30.56
40	ATOM	763	ND2	ASN	94	26.775	79.257	9.383	1.00	36.61
	ATOM	764	C	ASN	94	24.270	82.089	12.426	1.00	35.80
	ATOM	765	O	ASN	94	24.727	83.234	12.412	1.00	36.63
	ATOM	766	N	MET	95	23.649	81.585	13.484	1.00	35.73
	ATOM	767	CA	MET	95	23.476	82.375	14.697	1.00	34.73
45	ATOM	768	CB	MET	95	21.999	82.383	15.099	1.00	32.97
	ATOM	769	CG	MET	95	21.037	82.838	14.011	1.00	27.35
	ATOM	770	SD	MET	95	21.428	84.457	13.317	1.00	25.14
	ATOM	771	CE	MET	95	20.974	85.557	14.616	1.00	15.55
	ATOM	772	C	MET	95	24.321	81.856	15.862	1.00	36.36
50	ATOM	773	O	MET	95	24.642	80.665	15.924	1.00	40.70
	ATOM	774	N	PRO	96	24.717	82.749	16.784	1.00	36.06
	ATOM	775	CD	PRO	96	24.723	84.202	16.610	1.00	35.32
	ATOM	776	CA	PRO	96	25.516	82.388	17.956	1.00	40.00
	ATOM	777	CB	PRO	96	26.223	83.689	18.323	1.00	33.97
55	ATOM	778	CG	PRO	96	26.080	84.538	17.124	1.00	37.05
	ATOM	779	C	PRO	96	24.538	81.975	19.053	1.00	46.34
	ATOM	780	O	PRO	96	23.335	82.206	18.931	1.00	47.59
	ATOM	781	N	GLU	97	25.043	81.382	20.130	1.00	52.14
	ATOM	782	CA	GLU	97	24.177	80.956	21.223	1.00	57.27
60	ATOM	783	CB	GLU	97	24.814	79.795	21.987	1.00	62.29
	ATOM	784	CG	GLU	97	23.825	78.921	22.752	1.00	72.31
	ATOM	785	CD	GLU	97	23.481	77.631	21.992	1.00	79.34
	ATOM	786	OE1	GLU	97	24.416	76.806	21.747	1.00	84.94
	ATOM	787	OE2	GLU	97	22.284	77.445	21.639	1.00	81.39
65	ATOM	788	C	GLU	97	23.968	82.126	22.178	1.00	56.67
	ATOM	789	O	GLU	97	22.882	82.303	22.737	1.00	59.18

	ATOM	790	N	THR	98	25.009	82.933	22.344	1.00	57.55
	ATOM	791	CA	THR	98	24.974	84.068	23.259	1.00	59.71
	ATOM	792	CB	THR	98	26.372	84.642	23.449	1.00	59.24
	ATOM	793	OG1	THR	98	26.880	85.045	22.162	1.00	68.81
5	ATOM	794	CG2	THR	98	27.295	83.588	24.095	1.00	56.15
	ATOM	795	C	THR	98	24.045	85.227	22.900	1.00	60.32
	ATOM	796	O	THR	98	22.851	85.185	23.220	1.00	65.03
	ATOM	797	N	LEU	99	24.588	86.264	22.260	1.00	54.94
	ATOM	798	CA	LEU	99	23.794	87.441	21.898	1.00	51.13
10	ATOM	799	CB	LEU	99	24.517	88.706	22.358	1.00	50.56
	ATOM	800	CG	LEU	99	24.519	88.983	23.858	1.00	52.80
	ATOM	801	CD1	LEU	99	25.608	89.996	24.208	1.00	48.38
	ATOM	802	CD2	LEU	99	23.145	89.491	24.271	1.00	50.80
	ATOM	803	C	LEU	99	23.480	87.547	20.402	1.00	50.20
15	ATOM	804	O	LEU	99	24.099	88.341	19.679	1.00	50.50
	ATOM	805	N	PRO	100	22.487	86.776	19.925	1.00	47.97
	ATOM	806	CD	PRO	100	21.552	85.949	20.704	1.00	45.68
	ATOM	807	CA	PRO	100	22.104	86.785	18.511	1.00	46.54
	ATOM	808	CB	PRO	100	20.859	85.896	18.479	1.00	45.50
20	ATOM	809	CG	PRO	100	21.043	84.999	19.659	1.00	45.80
	ATOM	810	C	PRO	100	21.812	88.180	17.985	1.00	46.78
	ATOM	811	O	PRO	100	20.962	88.887	18.513	1.00	52.13
	ATOM	812	N	ASN	101	22.523	88.572	16.941	1.00	42.71
	ATOM	813	CA	ASN	101	22.328	89.867	16.318	1.00	41.37
25	ATOM	814	CB	ASN	101	22.928	90.970	17.181	1.00	43.24
	ATOM	815	CG	ASN	101	21.989	91.431	18.277	1.00	45.93
	ATOM	816	OD1	ASN	101	20.967	92.067	18.009	1.00	45.33
	ATOM	817	ND2	ASN	101	22.326	91.108	19.524	1.00	48.11
	ATOM	818	C	ASN	101	23.003	89.842	14.961	1.00	42.66
30	ATOM	819	O	ASN	101	24.135	90.304	14.811	1.00	44.79
	ATOM	820	N	ASN	102	22.319	89.293	13.961	1.00	40.46
	ATOM	821	CA	ASN	102	22.925	89.241	12.652	1.00	37.90
	ATOM	822	CB	ASN	102	23.043	87.802	12.175	1.00	34.01
	ATOM	823	CG	ASN	102	24.323	87.147	12.656	1.00	32.44
35	ATOM	824	OD1	ASN	102	25.380	87.772	12.666	1.00	29.07
	ATOM	825	ND2	ASN	102	24.236	85.885	13.050	1.00	33.79
	ATOM	826	C	ASN	102	22.365	90.110	11.543	1.00	39.20
	ATOM	827	O	ASN	102	23.058	91.000	11.064	1.00	46.07
	ATOM	828	N	SER	103	21.135	89.896	11.118	1.00	35.42
40	ATOM	829	CA	SER	103	20.620	90.721	10.012	1.00	40.28
	ATOM	830	CB	SER	103	20.771	92.235	10.303	1.00	40.68
	ATOM	831	OG	SER	103	21.868	92.823	9.613	1.00	31.76
	ATOM	832	C	SER	103	21.339	90.368	8.688	1.00	35.24
	ATOM	833	O	SER	103	22.561	90.440	8.571	1.00	24.65
45	ATOM	834	N	CYS	104	20.548	89.955	7.706	1.00	32.88
	ATOM	835	CA	CYS	104	21.064	89.583	6.414	1.00	31.77
	ATOM	836	C	CYS	104	20.312	90.400	5.347	1.00	30.58
	ATOM	837	O	CYS	104	19.096	90.552	5.409	1.00	29.23
	ATOM	838	CB	CYS	104	20.892	88.057	6.177	1.00	28.43
50	ATOM	839	SG	CYS	104	21.186	87.728	4.431	1.00	46.07
	ATOM	840	N	TYR	105	21.053	90.953	4.388	1.00	30.34
	ATOM	841	CA	TYR	105	20.473	91.721	3.288	1.00	26.12
	ATOM	842	CB	TYR	105	21.189	93.061	3.132	1.00	27.03
	ATOM	843	CG	TYR	105	20.794	93.857	1.897	1.00	25.61
55	ATOM	844	CD1	TYR	105	19.847	94.884	1.966	1.00	24.90
	ATOM	845	CE1	TYR	105	19.485	95.608	0.840	1.00	26.08
	ATOM	846	CD2	TYR	105	21.362	93.581	0.658	1.00	25.41
	ATOM	847	CE2	TYR	105	21.002	94.300	-0.474	1.00	28.91
	ATOM	848	CZ	TYR	105	20.065	95.312	-0.376	1.00	27.58
60	ATOM	849	OH	TYR	105	19.720	96.032	-1.499	1.00	29.89
	ATOM	850	C	TYR	105	20.645	90.916	2.002	1.00	29.20
	ATOM	851	O	TYR	105	21.649	90.236	1.815	1.00	33.10
	ATOM	852	N	SER	106	19.667	90.987	1.115	1.00	30.48
	ATOM	853	CA	SER	106	19.759	90.260	-0.141	1.00	29.22
65	ATOM	854	CB	SER	106	19.320	88.807	0.060	1.00	28.30
	ATOM	855	OG	SER	106	19.445	88.054	-1.130	1.00	28.62

	ATOM	856	C	SER	106	18.862	90.952	-1.152	1.00	29.94
	ATOM	857	O	SER	106	17.799	91.443	-0.792	1.00	29.72
	ATOM	858	N	ALA	107	19.299	91.011	-2.407	1.00	27.64
	ATOM	859	CA	ALA	107	18.518	91.650	-3.458	1.00	24.33
5	ATOM	860	CB	ALA	107	18.770	93.147	-3.460	1.00	21.30
	ATOM	861	C	ALA	107	18.832	91.070	-4.828	1.00	25.56
	ATOM	862	O	ALA	107	19.820	90.372	-5.012	1.00	28.03
	ATOM	863	N	GLY	108	17.978	91.369	-5.793	1.00	26.07
	ATOM	864	CA	GLY	108	18.187	90.882	-7.141	1.00	25.49
10	ATOM	865	C	GLY	108	17.130	91.436	-8.068	1.00	29.03
	ATOM	866	O	GLY	108	16.223	92.127	-7.624	1.00	32.20
	ATOM	867	N	ILE	109	17.240	91.138	-9.355	1.00	29.34
	ATOM	868	CA	ILE	109	16.278	91.618	-10.335	1.00	27.71
	ATOM	869	CB	ILE	109	16.992	92.217	-11.557	1.00	27.48
15	ATOM	870	CG2	ILE	109	15.979	92.693	-12.577	1.00	26.46
	ATOM	871	CG1	ILE	109	17.880	93.377	-11.109	1.00	26.16
	ATOM	872	CD1	ILE	109	18.755	93.934	-12.195	1.00	29.07
	ATOM	873	C	ILE	109	15.395	90.472	-10.790	1.00	28.66
	ATOM	874	O	ILE	109	15.857	89.341	-10.934	1.00	30.59
20	ATOM	875	N	ALA	110	14.120	90.759	-11.012	1.00	25.60
	ATOM	876	CA	ALA	110	13.174	89.743	-11.459	1.00	26.36
	ATOM	877	CB	ALA	110	12.517	89.087	-10.268	1.00	20.26
	ATOM	878	C	ALA	110	12.114	90.376	-12.342	1.00	29.61
	ATOM	879	O	ALA	110	11.845	91.563	-12.238	1.00	27.61
25	ATOM	880	N	LYS	111	11.522	89.593	-13.232	1.00	33.77
	ATOM	881	CA	LYS	111	10.480	90.132	-14.073	1.00	35.09
	ATOM	882	CB	LYS	111	10.532	89.544	-15.478	1.00	39.38
	ATOM	883	CG	LYS	111	9.333	89.997	-16.302	1.00	49.74
	ATOM	884	CD	LYS	111	9.654	90.191	-17.759	1.00	53.95
30	ATOM	885	CE	LYS	111	8.462	90.806	-18.477	1.00	54.28
	ATOM	886	NZ	LYS	111	8.754	90.983	-19.940	1.00	60.85
	ATOM	887	C	LYS	111	9.138	89.819	-13.423	1.00	35.28
	ATOM	888	O	LYS	111	8.847	88.673	-13.103	1.00	36.58
	ATOM	889	N	LEU	112	8.334	90.855	-13.221	1.00	32.32
35	ATOM	890	CA	LEU	112	7.034	90.710	-12.594	1.00	29.67
	ATOM	891	CB	LEU	112	6.989	91.527	-11.305	1.00	27.04
	ATOM	892	CG	LEU	112	8.154	91.327	-10.337	1.00	24.38
	ATOM	893	CD1	LEU	112	8.035	92.302	-9.183	1.00	27.56
	ATOM	894	CD2	LEU	112	8.167	89.908	-9.839	1.00	19.52
40	ATOM	895	C	LEU	112	5.940	91.188	-13.536	1.00	32.58
	ATOM	896	O	LEU	112	6.215	91.864	-14.527	1.00	31.51
	ATOM	897	N	GLU	113	4.698	90.841	-13.220	1.00	35.75
	ATOM	898	CA	GLU	113	3.569	91.235	-14.043	1.00	40.08
	ATOM	899	CB	GLU	113	3.006	90.044	-14.797	1.00	44.23
45	ATOM	900	CG	GLU	113	4.012	89.176	-15.493	1.00	50.12
	ATOM	901	CD	GLU	113	3.355	88.376	-16.594	1.00	58.08
	ATOM	902	OE1	GLU	113	2.192	87.919	-16.389	1.00	61.13
	ATOM	903	OE2	GLU	113	4.001	88.210	-17.661	1.00	63.64
	ATOM	904	C	GLU	113	2.442	91.806	-13.219	1.00	41.63
50	ATOM	905	O	GLU	113	2.361	91.575	-12.020	1.00	48.29
	ATOM	906	N	GLU	114	1.564	92.547	-13.888	1.00	40.97
	ATOM	907	CA	GLU	114	0.391	93.137	-13.257	1.00	36.68
	ATOM	908	CB	GLU	114	-0.592	93.601	-14.311	1.00	41.19
	ATOM	909	CG	GLU	114	-0.536	95.038	-14.689	1.00	46.58
55	ATOM	910	CD	GLU	114	-1.882	95.497	-15.205	1.00	51.12
	ATOM	911	OE1	GLU	114	-2.840	95.530	-14.394	1.00	51.88
	ATOM	912	OE2	GLU	114	-1.985	95.808	-16.417	1.00	57.08
	ATOM	913	C	GLU	114	-0.321	92.067	-12.456	1.00	35.51
	ATOM	914	O	GLU	114	-0.707	91.043	-13.006	1.00	38.92
60	ATOM	915	N	GLY	115	-0.512	92.301	-11.168	1.00	32.99
	ATOM	916	CA	GLY	115	-1.212	91.317	-10.373	1.00	31.79
	ATOM	917	C	GLY	115	-0.311	90.454	-9.530	1.00	34.39
	ATOM	918	O	GLY	115	-0.782	89.807	-8.605	1.00	39.20
	ATOM	919	N	ASP	116	0.973	90.417	-9.859	1.00	35.66
65	ATOM	920	CA	ASP	116	1.910	89.632	-9.074	1.00	38.09
	ATOM	921	CB	ASP	116	3.297	89.601	-9.728	1.00	36.00

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	ATOM	922	CG	ASP	116	3.376	88.654	-10.904	1.00	35.09
	ATOM	923	OD1	ASP	116	2.479	87.800	-11.061	1.00	33.81
	ATOM	924	OD2	ASP	116	4.354	88.752	-11.668	1.00	35.24
	ATOM	925	C	ASP	116	2.027	90.295	-7.709	1.00	39.38
5	ATOM	926	O	ASP	116	1.815	91.504	-7.575	1.00	37.91
	ATOM	927	N	GLU	117	2.350	89.506	-6.693	1.00	41.54
	ATOM	928	CA	GLU	117	2.527	90.052	-5.354	1.00	37.92
	ATOM	929	CB	GLU	117	1.433	89.558	-4.414	1.00	38.90
	ATOM	930	CG	GLU	117	0.037	89.897	-4.867	1.00	48.15
10	ATOM	931	CD	GLU	117	-1.012	89.424	-3.882	1.00	52.62
	ATOM	932	OE1	GLU	117	-0.837	88.320	-3.298	1.00	55.49
	ATOM	933	OE2	GLU	117	-2.016	90.157	-3.703	1.00	57.10
	ATOM	934	C	GLU	117	3.877	89.596	-4.836	1.00	34.02
	ATOM	935	O	GLU	117	4.316	88.487	-5.135	1.00	33.31
15	ATOM	936	N	LEU	118	4.544	90.463	-4.087	1.00	30.68
	ATOM	937	CA	LEU	118	5.833	90.134	-3.503	1.00	27.14
	ATOM	938	CB	LEU	118	6.861	91.219	-3.823	1.00	26.49
	ATOM	939	CG	LEU	118	7.276	91.421	-5.276	1.00	26.21
	ATOM	940	CD1	LEU	118	8.226	92.592	-5.368	1.00	23.05
20	ATOM	941	CD2	LEU	118	7.930	90.162	-5.808	1.00	25.38
	ATOM	942	C	LEU	118	5.665	90.045	-1.992	1.00	28.33
	ATOM	943	O	LEU	118	4.871	90.786	-1.408	1.00	31.46
	ATOM	944	N	GLN	119	6.394	89.135	-1.354	1.00	28.96
	ATOM	945	CA	GLN	119	6.323	89.015	0.096	1.00	29.13
25	ATOM	946	CB	GLN	119	5.202	88.066	0.494	1.00	28.23
	ATOM	947	CG	GLN	119	5.467	86.630	0.159	1.00	36.38
	ATOM	948	CD	GLN	119	4.301	85.737	0.510	1.00	36.57
	ATOM	949	OE1	GLN	119	4.446	84.517	0.573	1.00	40.70
	ATOM	950	NE2	GLN	119	3.136	86.335	0.730	1.00	36.80
30	ATOM	951	C	GLN	119	7.646	88.551	0.696	1.00	29.43
	ATOM	952	O	GLN	119	8.443	87.889	0.035	1.00	28.80
	ATOM	953	N	LEU	120	7.870	88.927	1.951	1.00	32.26
	ATOM	954	CA	LEU	120	9.077	88.584	2.697	1.00	29.96
	ATOM	955	CB	LEU	120	9.641	89.847	3.365	1.00	31.17
35	ATOM	956	CG	LEU	120	11.028	89.960	4.009	1.00	30.26
	ATOM	957	CD1	LEU	120	11.310	88.786	4.913	1.00	27.10
	ATOM	958	CD2	LEU	120	12.060	90.059	2.921	1.00	30.96
	ATOM	959	C	LEU	120	8.661	87.570	3.764	1.00	30.51
	ATOM	960	O	LEU	120	7.910	87.894	4.686	1.00	32.88
40	ATOM	961	N	ALA	121	9.154	86.343	3.640	1.00	27.40
	ATOM	962	CA	ALA	121	8.809	85.287	4.583	1.00	25.55
	ATOM	963	CB	ALA	121	8.059	84.186	3.852	1.00	19.41
	ATOM	964	C	ALA	121	10.005	84.690	5.326	1.00	29.89
	ATOM	965	O	ALA	121	11.063	84.461	4.745	1.00	35.88
45	ATOM	966	N	ILE	122	9.822	84.448	6.621	1.00	30.33
	ATOM	967	CA	ILE	122	10.859	83.846	7.448	1.00	30.27
	ATOM	968	CB	ILE	122	11.045	84.607	8.764	1.00	29.27
	ATOM	969	CG2	ILE	122	12.190	83.998	9.539	1.00	30.38
	ATOM	970	CG1	ILE	122	11.328	86.084	8.476	1.00	26.03
50	ATOM	971	CD1	ILE	122	11.498	86.949	9.711	1.00	29.21
	ATOM	972	C	ILE	122	10.380	82.429	7.749	1.00	33.20
	ATOM	973	O	ILE	122	9.369	82.242	8.421	1.00	33.30
	ATOM	974	N	PRO	123	11.101	81.412	7.245	1.00	34.70
	ATOM	975	CD	PRO	123	12.292	81.551	6.401	1.00	34.40
55	ATOM	976	CA	PRO	123	10.777	79.991	7.430	1.00	36.49
	ATOM	977	CB	PRO	123	11.673	79.286	6.407	1.00	30.22
	ATOM	978	CG	PRO	123	12.133	80.381	5.489	1.00	36.72
	ATOM	979	C	PRO	123	11.041	79.465	8.840	1.00	39.64
	ATOM	980	O	PRO	123	11.794	78.501	9.010	1.00	40.48
60	ATOM	981	N	ARG	124	10.426	80.091	9.840	1.00	44.58
	ATOM	982	CA	ARG	124	10.603	79.677	11.224	1.00	48.04
	ATOM	983	CB	ARG	124	11.794	80.398	11.844	1.00	52.91
	ATOM	984	CG	ARG	124	13.085	80.233	11.064	1.00	60.50
	ATOM	985	CD	ARG	124	14.019	79.334	11.843	1.00	72.57
65	ATOM	986	NE	ARG	124	15.229	78.994	11.074	1.00	84.90
	ATOM	987	CZ	ARG	124	15.367	77.925	10.279	1.00	85.44

	ATOM	988	NH1	ARG	124	14.357	77.065	10.132	1.00	87.69
	ATOM	989	NH2	ARG	124	16.517	77.719	9.626	1.00	82.78
	ATOM	990	C	ARG	124	9.356	79.935	12.054	1.00	48.39
	ATOM	991	O	ARG	124	8.625	80.895	11.806	1.00	45.26
5	ATOM	992	N	GLU	125	9.139	79.070	13.041	1.00	52.84
	ATOM	993	CA	GLU	125	7.982	79.135	13.931	1.00	55.02
	ATOM	994	CB	GLU	125	8.140	78.103	15.052	1.00	63.62
	ATOM	995	CG	GLU	125	9.594	77.803	15.423	1.00	70.62
	ATOM	996	CD	GLU	125	10.427	77.365	14.224	1.00	74.76
10	ATOM	997	OE1	GLU	125	10.137	76.277	13.656	1.00	75.75
	ATOM	998	OE2	GLU	125	11.366	78.111	13.850	1.00	80.71
	ATOM	999	C	GLU	125	7.697	80.506	14.528	1.00	53.44
	ATOM	1000	O	GLU	125	6.703	81.139	14.172	1.00	57.50
	ATOM	1001	N	ASN	126	8.524	80.955	15.461	1.00	47.18
15	ATOM	1002	CA	ASN	126	8.313	82.277	16.036	1.00	47.59
	ATOM	1003	CB	ASN	126	7.793	82.198	17.469	1.00	51.03
	ATOM	1004	CG	ASN	126	6.294	82.446	17.554	1.00	57.07
	ATOM	1005	OD1	ASN	126	5.477	81.577	17.212	1.00	61.15
	ATOM	1006	ND2	ASN	126	5.923	83.649	17.993	1.00	56.63
20	ATOM	1007	C	ASN	126	9.628	83.007	15.998	1.00	47.06
	ATOM	1008	O	ASN	126	10.267	83.249	17.026	1.00	45.61
	ATOM	1009	N	ALA	127	10.026	83.346	14.779	1.00	43.50
	ATOM	1010	CA	ALA	127	11.278	84.025	14.534	1.00	38.26
	ATOM	1011	CB	ALA	127	11.313	84.516	13.106	1.00	37.97
25	ATOM	1012	C	ALA	127	11.506	85.182	15.486	1.00	37.16
	ATOM	1013	O	ALA	127	10.646	86.046	15.652	1.00	33.14
	ATOM	1014	N	GLN	128	12.666	85.177	16.131	1.00	37.87
	ATOM	1015	CA	GLN	128	13.034	86.264	17.027	1.00	37.19
	ATOM	1016	CB	GLN	128	13.989	85.750	18.099	1.00	40.23
30	ATOM	1017	CG	GLN	128	13.309	84.803	19.059	1.00	38.14
	ATOM	1018	CD	GLN	128	12.076	85.431	19.664	1.00	39.51
	ATOM	1019	OE1	GLN	128	12.167	86.384	20.446	1.00	40.36
	ATOM	1020	NE2	GLN	128	10.905	84.919	19.289	1.00	41.53
	ATOM	1021	C	GLN	128	13.697	87.299	16.122	1.00	35.55
35	ATOM	1022	O	GLN	128	14.867	87.183	15.746	1.00	35.57
	ATOM	1023	N	ILE	129	12.912	88.307	15.775	1.00	34.64
	ATOM	1024	CA	ILE	129	13.313	89.355	14.847	1.00	29.60
	ATOM	1025	CB	ILE	129	12.401	89.229	13.573	1.00	31.03
	ATOM	1026	CG2	ILE	129	11.838	90.557	13.136	1.00	29.76
40	ATOM	1027	CG1	ILE	129	13.173	88.539	12.471	1.00	26.37
	ATOM	1028	CD1	ILE	129	13.598	87.168	12.841	1.00	34.70
	ATOM	1029	C	ILE	129	13.213	90.754	15.446	1.00	29.55
	ATOM	1030	O	ILE	129	12.443	90.986	16.366	1.00	33.42
	ATOM	1031	N	SER	130	14.004	91.689	14.931	1.00	27.67
45	ATOM	1032	CA	SER	130	13.934	93.069	15.402	1.00	25.00
	ATOM	1033	CB	SER	130	15.292	93.746	15.324	1.00	22.09
	ATOM	1034	OG	SER	130	15.152	95.140	15.558	1.00	20.89
	ATOM	1035	C	SER	130	12.967	93.817	14.498	1.00	26.03
	ATOM	1036	O	SER	130	13.018	93.660	13.291	1.00	30.77
50	ATOM	1037	N	LEU	131	12.094	94.631	15.070	1.00	24.49
	ATOM	1038	CA	LEU	131	11.128	95.365	14.267	1.00	28.07
	ATOM	1039	CB	LEU	131	9.717	95.164	14.821	1.00	28.87
	ATOM	1040	CG	LEU	131	8.944	93.956	14.295	1.00	30.76
	ATOM	1041	CD1	LEU	131	9.788	92.716	14.329	1.00	28.98
55	ATOM	1042	CD2	LEU	131	7.704	93.774	15.131	1.00	33.37
	ATOM	1043	C	LEU	131	11.412	96.851	14.129	1.00	33.66
	ATOM	1044	O	LEU	131	10.488	97.654	13.998	1.00	37.67
	ATOM	1045	N	ASP	132	12.689	97.218	14.160	1.00	34.97
	ATOM	1046	CA	ASP	132	13.078	98.616	14.002	1.00	34.31
60	ATOM	1047	CB	ASP	132	14.445	98.876	14.633	1.00	42.68
	ATOM	1048	CG	ASP	132	14.386	98.958	16.149	1.00	47.98
	ATOM	1049	OD1	ASP	132	15.467	98.958	16.788	1.00	50.41
	ATOM	1050	OD2	ASP	132	13.257	99.033	16.695	1.00	46.42
	ATOM	1051	C	ASP	132	13.120	98.935	12.517	1.00	34.32
65	ATOM	1052	O	ASP	132	13.676	98.181	11.727	1.00	32.87
	ATOM	1053	N	GLY	133	12.521	100.060	12.151	1.00	34.53

	ATOM	1054	CA	GLY	133	12.467	100.458	10.758	1.00	36.93
	ATOM	1055	C	GLY	133	13.780	100.538	10.006	1.00	35.06
	ATOM	1056	O	GLY	133	13.796	100.463	8.778	1.00	40.44
	ATOM	1057	N	ASP	134	14.886	100.684	10.721	1.00	31.08
5	ATOM	1058	CA	ASP	134	16.167	100.794	10.051	1.00	27.63
	ATOM	1059	CB	ASP	134	17.055	101.806	10.779	1.00	28.43
	ATOM	1060	CG	ASP	134	17.335	101.425	12.216	1.00	31.78
	ATOM	1061	OD1	ASP	134	16.453	100.829	12.861	1.00	35.68
	ATOM	1062	OD2	ASP	134	18.438	101.741	12.710	1.00	32.53
10	ATOM	1063	C	ASP	134	16.887	99.471	9.860	1.00	25.08
	ATOM	1064	O	ASP	134	17.728	99.356	8.997	1.00	19.71
	ATOM	1065	N	VAL	135	16.537	98.453	10.630	1.00	26.84
	ATOM	1066	CA	VAL	135	17.214	97.175	10.469	1.00	23.72
	ATOM	1067	CB	VAL	135	17.585	96.566	11.824	1.00	25.13
15	ATOM	1068	CG1	VAL	135	18.688	97.381	12.443	1.00	22.76
	ATOM	1069	CG2	VAL	135	16.375	96.522	12.731	1.00	22.12
	ATOM	1070	C	VAL	135	16.472	96.140	9.630	1.00	26.54
	ATOM	1071	O	VAL	135	17.099	95.255	9.063	1.00	29.26
	ATOM	1072	N	THR	136	15.145	96.227	9.549	1.00	26.39
20	ATOM	1073	CA	THR	136	14.414	95.280	8.710	1.00	28.40
	ATOM	1074	CB	THR	136	13.718	94.160	9.553	1.00	28.21
	ATOM	1075	OG1	THR	136	12.320	94.421	9.663	1.00	37.13
	ATOM	1076	CG2	THR	136	14.321	94.076	10.933	1.00	26.90
	ATOM	1077	C	THR	136	13.415	96.018	7.806	1.00	31.60
25	ATOM	1078	O	THR	136	12.485	96.678	8.277	1.00	27.29
	ATOM	1079	N	PHE	137	13.645	95.912	6.496	1.00	31.45
	ATOM	1080	CA	PHE	137	12.822	96.588	5.501	1.00	26.64
	ATOM	1081	CB	PHE	137	13.450	97.938	5.185	1.00	26.84
	ATOM	1082	CG	PHE	137	14.943	97.889	5.032	1.00	24.32
30	ATOM	1083	CD1	PHE	137	15.522	97.571	3.808	1.00	27.77
	ATOM	1084	CD2	PHE	137	15.771	98.169	6.106	1.00	23.70
	ATOM	1085	CE1	PHE	137	16.898	97.537	3.656	1.00	27.32
	ATOM	1086	CE2	PHE	137	17.150	98.134	5.962	1.00	26.31
	ATOM	1087	CZ	PHE	137	17.712	97.819	4.734	1.00	29.43
35	ATOM	1088	C	PHE	137	12.631	95.757	4.234	1.00	28.06
	ATOM	1089	O	PHE	137	13.344	94.788	4.021	1.00	25.04
	ATOM	1090	N	PHE	138	11.671	96.149	3.397	1.00	29.81
	ATOM	1091	CA	PHE	138	11.347	95.398	2.186	1.00	29.93
	ATOM	1092	CB	PHE	138	10.019	94.683	2.427	1.00	27.47
40	ATOM	1093	CG	PHE	138	9.683	93.648	1.408	1.00	28.49
	ATOM	1094	CD1	PHE	138	10.676	93.037	0.651	1.00	27.65
	ATOM	1095	CD2	PHE	138	8.356	93.275	1.207	1.00	31.06
	ATOM	1096	CE1	PHE	138	10.351	92.072	-0.293	1.00	30.90
	ATOM	1097	CE2	PHE	138	8.017	92.311	0.267	1.00	27.65
45	ATOM	1098	CZ	PHE	138	9.015	91.708	-0.486	1.00	27.41
	ATOM	1099	C	PHE	138	11.321	96.254	0.905	1.00	30.91
	ATOM	1100	O	PHE	138	10.652	97.279	0.831	1.00	24.94
	ATOM	1101	N	GLY	139	12.053	95.764	-0.100	1.00	36.94
	ATOM	1102	CA	GLY	139	12.284	96.413	-1.389	1.00	36.08
50	ATOM	1103	C	GLY	139	11.311	96.817	-2.470	1.00	36.48
	ATOM	1104	O	GLY	139	10.318	97.463	-2.190	1.00	43.61
	ATOM	1105	N	ALA	140	11.661	96.489	-3.715	1.00	36.60
	ATOM	1106	CA	ALA	140	10.873	96.790	-4.926	1.00	34.94
	ATOM	1107	CB	ALA	140	9.399	96.526	-4.675	1.00	36.94
55	ATOM	1108	C	ALA	140	11.039	98.176	-5.579	1.00	32.53
	ATOM	1109	O	ALA	140	10.520	99.178	-5.099	1.00	27.96
	ATOM	1110	N	LEU	141	11.752	98.201	-6.702	1.00	33.23
	ATOM	1111	CA	LEU	141	12.009	99.418	-7.483	1.00	36.11
	ATOM	1112	CB	LEU	141	13.411	99.942	-7.176	1.00	37.53
60	ATOM	1113	CG	LEU	141	13.983	101.155	-7.918	1.00	38.87
	ATOM	1114	CD1	LEU	141	15.253	101.589	-7.227	1.00	40.67
	ATOM	1115	CD2	LEU	141	14.285	100.828	-9.367	1.00	37.53
	ATOM	1116	C	LEU	141	11.902	99.068	-8.971	1.00	38.89
	ATOM	1117	O	LEU	141	12.527	98.114	-9.430	1.00	39.25
65	ATOM	1118	N	LYS	142	11.127	99.830	-9.735	1.00	42.86
	ATOM	1119	CA	LYS	142	10.986	99.514	-11.153	1.00	42.96

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	ATOM	1120	CB	LYS	142	9.632	99.990	-11.670	1.00	41.80
	ATOM	1121	CG	LYS	142	9.454	99.709	-13.143	1.00	46.88
	ATOM	1122	CD	LYS	142	8.001	99.724	-13.570	1.00	49.58
	ATOM	1123	CE	LYS	142	7.906	99.431	-15.058	1.00	51.07
5	ATOM	1124	NZ	LYS	142	6.503	99.348	-15.532	1.00	56.20
	ATOM	1125	C	LYS	142	12.094	100.058	-12.048	1.00	43.41
	ATOM	1126	O	LYS	142	12.422	101.240	-11.999	1.00	44.98
	ATOM	1127	N	LEU	143	12.664	99.183	-12.872	1.00	42.81
	ATOM	1128	CA	LEU	143	13.734	99.570	-13.789	1.00	40.65
10	ATOM	1129	CB	LEU	143	14.597	98.359	-14.138	1.00	36.56
	ATOM	1130	CG	LEU	143	15.264	97.589	-12.999	1.00	36.17
	ATOM	1131	CD1	LEU	143	16.013	96.401	-13.573	1.00	37.95
	ATOM	1132	CD2	LEU	143	16.208	98.497	-12.241	1.00	27.38
	ATOM	1133	C	LEU	143	13.151	100.136	-15.079	1.00	43.14
15	ATOM	1134	O	LEU	143	12.066	99.733	-15.509	1.00	46.28
	ATOM	1135	N	LEU	144	13.866	101.066	-15.699	1.00	42.63
	ATOM	1136	CA	LEU	144	13.396	101.648	-16.947	1.00	44.11
	ATOM	1137	CB	LEU	144	14.085	102.984	-17.209	1.00	44.88
	ATOM	1138	CG	LEU	144	13.752	104.076	-16.195	1.00	47.02
20	ATOM	1139	CD1	LEU	144	14.558	105.313	-16.487	1.00	47.71
	ATOM	1140	CD2	LEU	144	12.268	104.389	-16.247	1.00	47.87
	ATOM	1141	C	LEU	144	13.685	100.704	-18.097	1.00	45.66
	ATOM	1142	O	LEU	144	14.564	99.835	-17.942	1.00	46.37
	ATOM	1143	OXT	LEU	144	13.034	100.854	-19.148	1.00	50.66
25	END					-23.152	-5.163	-22.411	0.00	0.00

TABLE 9

11									
5	ATOM	1	CB	VAL	1	-32.869	106.897	-43.833	1.00 69.28
	ATOM	2	CG1	VAL	1	-33.490	108.285	-44.101	1.00 68.26
	ATOM	3	CG2	VAL	1	-31.354	106.913	-44.062	1.00 71.75
	ATOM	4	C	VAL	1	-34.681	106.050	-42.314	1.00 64.86
	ATOM	5	O	VAL	1	-35.505	106.801	-41.772	1.00 64.02
10	ATOM	6	N	VAL	1	-32.327	105.285	-41.966	1.00 65.10
	ATOM	7	CA	VAL	1	-33.190	106.435	-42.375	1.00 65.98
	ATOM	8	N	THR	2	-35.023	104.889	-42.879	1.00 61.42
	ATOM	9	CA	THR	2	-36.409	104.422	-42.876	1.00 56.64
	ATOM	10	CB	THR	2	-36.956	104.233	-44.302	1.00 56.77
15	ATOM	11	OG1	THR	2	-36.091	103.349	-45.021	1.00 57.45
	ATOM	12	CG2	THR	2	-37.047	105.566	-45.035	1.00 55.68
	ATOM	13	C	THR	2	-36.549	103.089	-42.168	1.00 53.08
	ATOM	14	O	THR	2	-35.562	102.430	-41.856	1.00 52.00
	ATOM	15	N	GLN	3	-37.792	102.688	-41.933	1.00 50.34
20	ATOM	16	CA	GLN	3	-38.073	101.425	-41.261	1.00 45.97
	ATOM	17	CB	GLN	3	-39.006	101.659	-40.079	1.00 46.09
	ATOM	18	CG	GLN	3	-38.577	102.786	-39.176	1.00 49.60
	ATOM	19	CD	GLN	3	-39.483	102.926	-37.980	1.00 50.43
	ATOM	20	OE1	GLN	3	-39.510	102.056	-37.115	1.00 50.94
25	ATOM	21	NE2	GLN	3	-40.239	104.020	-37.925	1.00 49.99
	ATOM	22	C	GLN	3	-38.715	100.435	-42.215	1.00 42.41
	ATOM	23	O	GLN	3	-39.863	100.595	-42.619	1.00 37.34
	ATOM	24	N	ASP	4	-37.965	99.407	-42.576	1.00 40.32
	ATOM	25	CA	ASP	4	-38.477	98.394	-43.482	1.00 39.37
30	ATOM	26	CB	ASP	4	-37.350	97.448	-43.905	1.00 44.56
	ATOM	27	CG	ASP	4	-36.288	98.142	-44.731	1.00 45.76
	ATOM	28	OD1	ASP	4	-36.220	99.392	-44.673	1.00 48.58
	ATOM	29	OD2	ASP	4	-35.520	97.443	-45.427	1.00 45.55
	ATOM	30	C	ASP	4	-39.563	97.607	-42.782	1.00 36.30
35	ATOM	31	O	ASP	4	-39.515	97.405	-41.571	1.00 36.29
	ATOM	32	N	CYS	5	-40.551	97.174	-43.548	1.00 35.11
	ATOM	33	CA	CYS	5	-41.634	96.376	-43.004	1.00 35.61
	ATOM	34	CB	CYS	5	-42.626	97.247	-42.228	1.00 34.96
	ATOM	35	SG	CYS	5	-43.038	98.801	-42.995	1.00 35.34
40	ATOM	36	C	CYS	5	-42.331	95.632	-44.124	1.00 31.80
	ATOM	37	O	CYS	5	-42.338	96.076	-45.266	1.00 31.85
	ATOM	38	N	LEU	6	-42.882	94.476	-43.792	1.00 27.72
	ATOM	39	CA	LEU	6	-43.592	93.658	-44.753	1.00 30.80
	ATOM	40	CB	LEU	6	-42.705	92.513	-45.228	1.00 29.25
45	ATOM	41	CG	LEU	6	-43.314	91.495	-46.194	1.00 29.47
	ATOM	42	CD1	LEU	6	-42.223	90.851	-47.006	1.00 30.65
	ATOM	43	CD2	LEU	6	-44.062	90.448	-45.424	1.00 35.27
	ATOM	44	C	LEU	6	-44.826	93.119	-44.056	1.00 31.83
	ATOM	45	O	LEU	6	-44.726	92.550	-42.975	1.00 33.45
50	ATOM	46	N	GLN	7	-45.991	93.307	-44.665	1.00 32.23
	ATOM	47	CA	GLN	7	-47.225	92.842	-44.069	1.00 30.26
	ATOM	48	CB	GLN	7	-48.145	94.027	-43.806	1.00 29.56
	ATOM	49	CG	GLN	7	-49.337	93.726	-42.924	1.00 25.09
	ATOM	50	CD	GLN	7	-50.053	94.992	-42.499	1.00 26.50
55	ATOM	51	OE1	GLN	7	-50.709	95.655	-43.302	1.00 24.40
	ATOM	52	NE2	GLN	7	-49.915	95.344	-41.233	1.00 27.03
	ATOM	53	C	GLN	7	-47.901	91.838	-44.978	1.00 32.11
	ATOM	54	O	GLN	7	-47.902	91.986	-46.193	1.00 34.24
	ATOM	55	N	LEU	8	-48.460	90.804	-44.367	1.00 33.73
60	ATOM	56	CA	LEU	8	-49.153	89.749	-45.088	1.00 35.41
	ATOM	57	CB	LEU	8	-48.511	88.387	-44.771	1.00 32.44
	ATOM	58	CG	LEU	8	-47.405	87.813	-45.669	1.00 27.69
	ATOM	59	CD1	LEU	8	-46.768	88.879	-46.522	1.00 26.43
	ATOM	60	CD2	LEU	8	-46.392	87.127	-44.793	1.00 18.64
65	ATOM	61	C	LEU	8	-50.627	89.738	-44.712	1.00 36.87
	ATOM	62	O	LEU	8	-51.002	90.051	-43.583	1.00 36.05
	ATOM	63	N	ILE	9	-51.456	89.363	-45.674	1.00 39.25

	ATOM	64	CA	ILE	9	-52.898	89.316	-45.497	1.00	35.57
	ATOM	65	CB	ILE	9	-53.541	90.413	-46.382	1.00	34.28
	ATOM	66	CG2	ILE	9	-54.730	89.889	-47.148	1.00	38.20
	ATOM	67	CG1	ILE	9	-53.933	91.591	-45.520	1.00	35.22
5	ATOM	68	CD1	ILE	9	-54.613	92.652	-46.318	1.00	43.63
	ATOM	69	C	ILE	9	-53.415	87.924	-45.872	1.00	35.28
	ATOM	70	O	ILE	9	-52.864	87.271	-46.754	1.00	33.75
	ATOM	71	N	ALA	10	-54.458	87.460	-45.194	1.00	35.78
	ATOM	72	CA	ALA	10	-55.008	86.149	-45.506	1.00	35.28
10	ATOM	73	CB	ALA	10	-56.108	85.797	-44.531	1.00	28.47
	ATOM	74	C	ALA	10	-55.548	86.122	-46.929	1.00	36.49
	ATOM	75	O	ALA	10	-56.171	87.075	-47.390	1.00	37.41
	ATOM	76	N	ASP	11	-55.302	85.022	-47.626	1.00	39.44
	ATOM	77	CA	ASP	11	-55.770	84.862	-48.997	1.00	40.77
15	ATOM	78	CB	ASP	11	-54.641	84.324	-49.876	1.00	38.90
	ATOM	79	CG	ASP	11	-55.101	84.014	-51.280	1.00	40.34
	ATOM	80	OD1	ASP	11	-56.093	84.629	-51.738	1.00	39.50
	ATOM	81	OD2	ASP	11	-54.461	83.162	-51.924	1.00	37.38
	ATOM	82	C	ASP	11	-56.966	83.917	-49.035	1.00	42.25
20	ATOM	83	O	ASP	11	-56.813	82.696	-49.108	1.00	40.54
	ATOM	84	N	SER	12	-58.157	84.500	-48.991	1.00	42.92
	ATOM	85	CA	SER	12	-59.406	83.742	-48.978	1.00	43.84
	ATOM	86	CB	SER	12	-60.579	84.695	-48.782	1.00	42.13
	ATOM	87	OG	SER	12	-60.602	85.670	-49.809	1.00	43.31
25	ATOM	88	C	SER	12	-59.663	82.896	-50.215	1.00	46.10
	ATOM	89	O	SER	12	-60.641	82.150	-50.268	1.00	43.90
	ATOM	90	N	GLU	13	-58.785	83.005	-51.205	1.00	46.72
	ATOM	91	CA	GLU	13	-58.960	82.247	-52.429	1.00	44.93
	ATOM	92	CB	GLU	13	-58.736	83.150	-53.629	1.00	46.71
30	ATOM	93	CG	GLU	13	-59.805	84.202	-53.739	1.00	58.25
	ATOM	94	CD	GLU	13	-59.970	84.704	-55.159	1.00	66.70
	ATOM	95	OE1	GLU	13	-60.252	83.855	-56.049	1.00	74.10
	ATOM	96	OE2	GLU	13	-59.822	85.935	-55.389	1.00	71.55
	ATOM	97	C	GLU	13	-58.130	80.974	-52.557	1.00	43.24
35	ATOM	98	O	GLU	13	-58.026	80.402	-53.644	1.00	44.25
	ATOM	99	N	THR	14	-57.529	80.534	-51.461	1.00	37.06
	ATOM	100	CA	THR	14	-56.781	79.283	-51.486	1.00	36.77
	ATOM	101	CB	THR	14	-55.251	79.490	-51.689	1.00	33.39
	ATOM	102	OG1	THR	14	-54.660	79.980	-50.487	1.00	41.69
40	ATOM	103	CG2	THR	14	-54.989	80.475	-52.817	1.00	31.66
	ATOM	104	C	THR	14	-57.054	78.593	-50.154	1.00	34.96
	ATOM	105	O	THR	14	-57.301	79.247	-49.147	1.00	35.04
	ATOM	106	N	PRO	15	-57.034	77.261	-50.140	1.00	37.11
	ATOM	107	CD	PRO	15	-56.780	76.385	-51.293	1.00	36.50
45	ATOM	108	CA	PRO	15	-57.288	76.471	-48.929	1.00	38.81
	ATOM	109	CB	PRO	15	-57.278	75.033	-49.446	1.00	39.15
	ATOM	110	CG	PRO	15	-57.569	75.173	-50.921	1.00	37.32
	ATOM	111	C	PRO	15	-56.231	76.677	-47.852	1.00	37.78
	ATOM	112	O	PRO	15	-55.083	76.981	-48.163	1.00	35.91
50	ATOM	113	N	THR	16	-56.608	76.510	-46.589	1.00	36.67
	ATOM	114	CA	THR	16	-55.633	76.666	-45.522	1.00	38.46
	ATOM	115	CB	THR	16	-56.285	76.614	-44.127	1.00	38.61
	ATOM	116	OG1	THR	16	-56.869	75.324	-43.916	1.00	42.44
	ATOM	117	CG2	THR	16	-57.361	77.681	-44.006	1.00	40.11
55	ATOM	118	C	THR	16	-54.686	75.493	-45.656	1.00	35.97
	ATOM	119	O	THR	16	-55.126	74.379	-45.909	1.00	39.85
	ATOM	120	N	ILE	17	-53.392	75.739	-45.496	1.00	33.66
	ATOM	121	CA	ILE	17	-52.403	74.678	-45.619	1.00	34.76
	ATOM	122	CB	ILE	17	-51.002	75.265	-45.832	1.00	32.44
60	ATOM	123	CG2	ILE	17	-49.967	74.157	-45.836	1.00	28.66
	ATOM	124	CG1	ILE	17	-50.976	76.059	-47.143	1.00	33.15
	ATOM	125	CD1	ILE	17	-49.662	76.758	-47.427	1.00	31.19
	ATOM	126	C	ILE	17	-52.364	73.743	-44.417	1.00	40.64
	ATOM	127	O	ILE	17	-52.247	74.184	-43.281	1.00	44.07
65	ATOM	128	N	GLN	18	-52.465	72.442	-44.673	1.00	45.91
	ATOM	129	CA	GLN	18	-52.426	71.444	-43.607	1.00	50.22

	ATOM	130	CB	GLN	18	-53.528	70.407	-43.796	1.00	53.09
	ATOM	131	CG	GLN	18	-54.443	70.307	-42.604	1.00	57.60
	ATOM	132	CD	GLN	18	-55.271	71.556	-42.452	1.00	61.39
	ATOM	133	OE1	GLN	18	-56.156	71.830	-43.267	1.00	68.36
5	ATOM	134	NE2	GLN	18	-54.981	72.336	-41.419	1.00	64.93
	ATOM	135	C	GLN	18	-51.087	70.729	-43.591	1.00	51.56
	ATOM	136	O	GLN	18	-50.597	70.287	-44.626	1.00	56.44
	ATOM	137	N	LYS	19	-50.495	70.607	-42.415	1.00	52.60
	ATOM	138	CA	LYS	19	-49.212	69.934	-42.312	1.00	56.10
10	ATOM	139	CB	LYS	19	-48.112	70.771	-42.967	1.00	55.55
	ATOM	140	CG	LYS	19	-46.747	70.150	-42.781	1.00	61.15
	ATOM	141	CD	LYS	19	-45.667	70.829	-43.599	1.00	64.42
	ATOM	142	CE	LYS	19	-44.317	70.133	-43.372	1.00	64.84
	ATOM	143	NZ	LYS	19	-43.210	70.735	-44.182	1.00	69.52
15	ATOM	144	C	LYS	19	-48.823	69.629	-40.873	1.00	57.41
	ATOM	145	O	LYS	19	-48.947	70.487	-39.996	1.00	61.02
	ATOM	146	N	GLY	20	-48.346	68.406	-40.644	1.00	58.84
	ATOM	147	CA	GLY	20	-47.936	67.991	-39.316	1.00	56.66
	ATOM	148	C	GLY	20	-48.968	68.305	-38.251	1.00	56.87
20	ATOM	149	O	GLY	20	-48.604	68.680	-37.135	1.00	58.64
	ATOM	150	N	SER	21	-50.249	68.151	-38.594	1.00	54.49
	ATOM	151	CA	SER	21	-51.355	68.420	-37.672	1.00	53.68
	ATOM	152	CB	SER	21	-51.270	67.497	-36.449	1.00	55.80
	ATOM	153	OG	SER	21	-50.411	68.021	-35.453	1.00	65.34
25	ATOM	154	C	SER	21	-51.411	69.901	-37.226	1.00	51.41
	ATOM	155	O	SER	21	-51.911	70.235	-36.138	1.00	47.74
	ATOM	156	N	TYR	22	-50.886	70.776	-38.082	1.00	46.25
	ATOM	157	CA	TYR	22	-50.879	72.214	-37.849	1.00	40.57
	ATOM	158	CB	TYR	22	-49.445	72.744	-37.787	1.00	43.27
30	ATOM	159	CG	TYR	22	-48.822	72.683	-36.416	1.00	49.18
	ATOM	160	CD1	TYR	22	-48.679	71.473	-35.746	1.00	52.61
	ATOM	161	CE1	TYR	22	-48.114	71.414	-34.469	1.00	51.92
	ATOM	162	CD2	TYR	22	-48.383	73.839	-35.779	1.00	50.01
	ATOM	163	CE2	TYR	22	-47.815	73.791	-34.505	1.00	50.94
35	ATOM	164	CZ	TYR	22	-47.685	72.576	-33.857	1.00	50.78
	ATOM	165	OH	TYR	22	-47.127	72.529	-32.599	1.00	53.87
	ATOM	166	C	TYR	22	-51.606	72.874	-39.015	1.00	37.85
	ATOM	167	O	TYR	22	-51.580	72.364	-40.135	1.00	38.70
	ATOM	168	N	THR	23	-52.271	73.993	-38.759	1.00	33.81
40	ATOM	169	CA	THR	23	-52.964	74.703	-39.828	1.00	32.40
	ATOM	170	CB	THR	23	-54.411	75.103	-39.441	1.00	31.58
	ATOM	171	OG1	THR	23	-55.158	73.943	-39.062	1.00	30.76
	ATOM	172	CG2	THR	23	-55.097	75.771	-40.613	1.00	24.08
	ATOM	173	C	THR	23	-52.190	75.977	-40.115	1.00	33.28
45	ATOM	174	O	THR	23	-51.833	76.708	-39.197	1.00	30.44
	ATOM	175	N	PHE	24	-51.928	76.237	-41.387	1.00	32.16
	ATOM	176	CA	PHE	24	-51.202	77.433	-41.778	1.00	30.68
	ATOM	177	CB	PHE	24	-49.905	77.063	-42.495	1.00	28.20
	ATOM	178	CG	PHE	24	-48.909	76.373	-41.618	1.00	31.00
50	ATOM	179	CD1	PHE	24	-48.979	75.006	-41.408	1.00	31.69
	ATOM	180	CD2	PHE	24	-47.912	77.099	-40.971	1.00	31.42
	ATOM	181	CE1	PHE	24	-48.074	74.372	-40.566	1.00	32.57
	ATOM	182	CE2	PHE	24	-47.007	76.473	-40.131	1.00	25.01
	ATOM	183	CZ	PHE	24	-47.088	75.107	-39.929	1.00	30.12
55	ATOM	184	C	PHE	24	-52.043	78.335	-42.672	1.00	31.73
	ATOM	185	O	PHE	24	-52.627	77.890	-43.655	1.00	33.31
	ATOM	186	N	VAL	25	-52.106	79.608	-42.315	1.00	32.92
	ATOM	187	CA	VAL	25	-52.861	80.571	-43.088	1.00	30.84
	ATOM	188	CB	VAL	25	-52.956	81.915	-42.348	1.00	30.29
60	ATOM	189	CG1	VAL	25	-53.710	82.916	-43.187	1.00	28.13
	ATOM	190	CG2	VAL	25	-53.631	81.723	-41.017	1.00	29.53
	ATOM	191	C	VAL	25	-52.176	80.818	-44.427	1.00	36.08
	ATOM	192	O	VAL	25	-50.956	80.948	-44.495	1.00	35.82
	ATOM	193	N	PRO	26	-52.954	80.859	-45.518	1.00	38.03
65	ATOM	194	CD	PRO	26	-54.355	80.416	-45.593	1.00	38.28
	ATOM	195	CA	PRO	26	-52.419	81.100	-46.865	1.00	38.08

	ATOM	196	CB	PRO	26	-53.546	80.622	-47.784	1.00	37.62
	ATOM	197	CG	PRO	26	-54.379	79.730	-46.920	1.00	36.31
	ATOM	198	C	PRO	26	-52.224	82.611	-46.976	1.00	40.85
	ATOM	199	O	PRO	26	-53.199	83.360	-47.000	1.00	42.27
5	ATOM	200	N	TRP	27	-50.984	83.072	-47.044	1.00	40.86
	ATOM	201	CA	TRP	27	-50.759	84.511	-47.104	1.00	37.05
	ATOM	202	CB	TRP	27	-49.460	84.878	-46.384	1.00	33.29
	ATOM	203	CG	TRP	27	-49.453	84.504	-44.946	1.00	31.06
	ATOM	204	CD2	TRP	27	-50.341	84.979	-43.933	1.00	27.79
10	ATOM	205	CE2	TRP	27	-49.987	84.334	-42.737	1.00	29.28
	ATOM	206	CE3	TRP	27	-51.402	85.885	-43.920	1.00	27.06
	ATOM	207	CD1	TRP	27	-48.620	83.623	-44.342	1.00	31.07
	ATOM	208	NE1	TRP	27	-48.930	83.511	-43.016	1.00	32.70
	ATOM	209	CZ2	TRP	27	-50.658	84.564	-41.532	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-52.070	86.114	-42.722	1.00	26.93
	ATOM	211	CH2	TRP	27	-51.693	85.454	-41.546	1.00	28.08
	ATOM	212	C	TRP	27	-50.741	85.117	-48.489	1.00	38.50
	ATOM	213	O	TRP	27	-50.509	84.438	-49.483	1.00	39.07
	ATOM	214	N	LEU	28	-50.999	86.419	-48.521	1.00	40.19
20	ATOM	215	CA	LEU	28	-51.002	87.210	-49.741	1.00	39.45
	ATOM	216	CB	LEU	28	-52.432	87.526	-50.163	1.00	44.18
	ATOM	217	CG	LEU	28	-52.609	88.004	-51.603	1.00	46.83
	ATOM	218	CD1	LEU	28	-52.276	86.849	-52.551	1.00	45.15
	ATOM	219	CD2	LEU	28	-54.047	88.487	-51.816	1.00	49.15
25	ATOM	220	C	LEU	28	-50.297	88.499	-49.343	1.00	39.80
	ATOM	221	O	LEU	28	-50.528	89.011	-48.257	1.00	42.40
	ATOM	222	N	LEU	29	-49.434	89.023	-50.197	1.00	33.58
	ATOM	223	CA	LEU	29	-48.738	90.245	-49.846	1.00	29.21
	ATOM	224	CB	LEU	29	-47.680	90.591	-50.893	1.00	27.16
30	ATOM	225	CG	LEU	29	-46.944	91.908	-50.628	1.00	22.36
	ATOM	226	CD1	LEU	29	-45.971	91.726	-49.475	1.00	21.47
	ATOM	227	CD2	LEU	29	-46.219	92.362	-51.867	1.00	14.53
	ATOM	228	C	LEU	29	-49.681	91.433	-49.696	1.00	29.98
	ATOM	229	O	LEU	29	-50.471	91.737	-50.583	1.00	30.36
35	ATOM	230	N	SER	30	-49.588	92.097	-48.551	1.00	29.88
	ATOM	231	CA	SER	30	-50.391	93.276	-48.288	1.00	28.49
	ATOM	232	CB	SER	30	-50.647	93.436	-46.789	1.00	26.13
	ATOM	233	OG	SER	30	-51.347	94.632	-46.528	1.00	20.25
	ATOM	234	C	SER	30	-49.545	94.430	-48.809	1.00	31.61
40	ATOM	235	O	SER	30	-50.002	95.238	-49.612	1.00	35.14
	ATOM	236	N	PHE	31	-48.299	94.493	-48.351	1.00	32.12
	ATOM	237	CA	PHE	31	-47.375	95.523	-48.793	1.00	30.76
	ATOM	238	CB	PHE	31	-47.815	96.905	-48.293	1.00	27.55
	ATOM	239	CG	PHE	31	-47.290	97.266	-46.931	1.00	31.23
45	ATOM	240	CD1	PHE	31	-46.009	97.785	-46.780	1.00	32.00
	ATOM	241	CD2	PHE	31	-48.072	97.076	-45.794	1.00	27.89
	ATOM	242	CE1	PHE	31	-45.517	98.107	-45.525	1.00	29.70
	ATOM	243	CE2	PHE	31	-47.585	97.397	-44.538	1.00	26.15
	ATOM	244	CZ	PHE	31	-46.307	97.912	-44.402	1.00	28.30
50	ATOM	245	C	PHE	31	-45.974	95.205	-48.301	1.00	32.06
	ATOM	246	O	PHE	31	-45.790	94.565	-47.266	1.00	32.26
	ATOM	247	N	LYS	32	-44.987	95.645	-49.063	1.00	34.83
	ATOM	248	CA	LYS	32	-43.600	95.431	-48.705	1.00	37.13
	ATOM	249	CB	LYS	32	-42.961	94.421	-49.657	1.00	36.55
55	ATOM	250	CG	LYS	32	-41.499	94.193	-49.392	1.00	40.62
	ATOM	251	CD	LYS	32	-40.836	93.421	-50.505	1.00	44.56
	ATOM	252	CE	LYS	32	-39.317	93.490	-50.330	1.00	46.29
	ATOM	253	NZ	LYS	32	-38.575	92.649	-51.316	1.00	50.58
	ATOM	254	C	LYS	32	-42.903	96.784	-48.811	1.00	36.39
60	ATOM	255	O	LYS	32	-43.089	97.510	-49.788	1.00	38.69
	ATOM	256	N	ARG	33	-42.118	97.132	-47.798	1.00	37.90
	ATOM	257	CA	ARG	33	-41.413	98.406	-47.796	1.00	35.74
	ATOM	258	CB	ARG	33	-42.101	99.357	-46.823	1.00	36.79
	ATOM	259	CG	ARG	33	-41.437	100.704	-46.649	1.00	31.09
65	ATOM	260	CD	ARG	33	-42.365	101.636	-45.880	1.00	31.62
	ATOM	261	NE	ARG	33	-41.870	103.007	-45.795	1.00	38.16

	ATOM	262	CZ	ARG	33	-41.269	103.519	-44.727	1.00	39.15
	ATOM	263	NH1	ARG	33	-41.089	102.775	-43.645	1.00	43.41
	ATOM	264	NH2	ARG	33	-40.845	104.772	-44.738	1.00	40.65
	ATOM	265	C	ARG	33	-39.956	98.199	-47.398	1.00	37.71
5	ATOM	266	O	ARG	33	-39.663	97.720	-46.299	1.00	41.16
	ATOM	267	N	GLY	34	-39.037	98.548	-48.291	1.00	36.59
	ATOM	268	CA	GLY	34	-37.634	98.377	-47.977	1.00	36.72
	ATOM	269	C	GLY	34	-37.087	97.046	-48.447	1.00	40.01
	ATOM	270	O	GLY	34	-37.716	96.341	-49.236	1.00	39.85
10	ATOM	271	N	SER	35	-35.917	96.690	-47.934	1.00	38.74
	ATOM	272	CA	SER	35	-35.252	95.456	-48.326	1.00	40.23
	ATOM	273	CB	SER	35	-33.905	95.806	-48.944	1.00	43.02
	ATOM	274	OG	SER	35	-33.152	96.631	-48.055	1.00	48.05
	ATOM	275	C	SER	35	-35.025	94.433	-47.210	1.00	40.54
15	ATOM	276	O	SER	35	-34.770	93.264	-47.495	1.00	43.74
	ATOM	277	N	ALA	36	-35.116	94.864	-45.954	1.00	35.56
	ATOM	278	CA	ALA	36	-34.876	93.985	-44.815	1.00	29.62
	ATOM	279	CB	ALA	36	-34.957	94.789	-43.534	1.00	25.47
	ATOM	280	C	ALA	36	-35.768	92.752	-44.706	1.00	31.23
20	ATOM	281	O	ALA	36	-35.372	91.761	-44.105	1.00	29.56
	ATOM	282	N	LEU	37	-36.964	92.802	-45.282	1.00	34.09
	ATOM	283	CA	LEU	37	-37.891	91.677	-45.197	1.00	32.32
	ATOM	284	CB	LEU	37	-39.035	92.022	-44.246	1.00	26.47
	ATOM	285	CG	LEU	37	-38.607	92.350	-42.818	1.00	25.72
25	ATOM	286	CD1	LEU	37	-39.697	93.111	-42.105	1.00	24.61
	ATOM	287	CD2	LEU	37	-38.262	91.075	-42.086	1.00	27.53
	ATOM	288	C	LEU	37	-38.459	91.270	-46.545	1.00	35.16
	ATOM	289	O	LEU	37	-38.746	92.108	-47.392	1.00	36.47
	ATOM	290	N	GLU	38	-38.625	89.969	-46.729	1.00	38.30
30	ATOM	291	CA	GLU	38	-39.163	89.427	-47.967	1.00	41.17
	ATOM	292	CB	GLU	38	-38.044	88.845	-48.822	1.00	46.15
	ATOM	293	CG	GLU	38	-37.374	89.814	-49.767	1.00	50.01
	ATOM	294	CD	GLU	38	-36.103	89.225	-50.365	1.00	51.18
	ATOM	295	OE1	GLU	38	-36.112	88.010	-50.692	1.00	48.40
35	ATOM	296	OE2	GLU	38	-35.102	89.976	-50.505	1.00	50.23
	ATOM	297	C	GLU	38	-40.157	88.316	-47.689	1.00	42.69
	ATOM	298	O	GLU	38	-40.131	87.691	-46.637	1.00	42.55
	ATOM	299	N	GLU	39	-41.039	88.066	-48.642	1.00	44.75
	ATOM	300	CA	GLU	39	-42.002	86.989	-48.478	1.00	46.50
40	ATOM	301	CB	GLU	39	-43.330	87.352	-49.135	1.00	47.52
	ATOM	302	CG	GLU	39	-43.157	88.097	-50.441	1.00	58.50
	ATOM	303	CD	GLU	39	-44.474	88.331	-51.156	1.00	62.35
	ATOM	304	OE1	GLU	39	-44.488	89.106	-52.147	1.00	64.58
	ATOM	305	OE2	GLU	39	-45.490	87.731	-50.732	1.00	63.04
45	ATOM	306	C	GLU	39	-41.361	85.812	-49.189	1.00	44.56
	ATOM	307	O	GLU	39	-40.789	85.971	-50.260	1.00	48.11
	ATOM	308	N	LYS	40	-41.433	84.635	-48.588	1.00	40.71
	ATOM	309	CA	LYS	40	-40.837	83.459	-49.189	1.00	38.08
	ATOM	310	CB	LYS	40	-39.371	83.342	-48.776	1.00	41.28
50	ATOM	311	CG	LYS	40	-38.705	82.043	-49.210	1.00	41.52
	ATOM	312	CD	LYS	40	-37.377	81.845	-48.493	1.00	42.73
	ATOM	313	CE	LYS	40	-36.765	80.486	-48.783	1.00	41.97
	ATOM	314	NZ	LYS	40	-35.528	80.274	-47.970	1.00	45.94
	ATOM	315	C	LYS	40	-41.573	82.202	-48.776	1.00	39.66
55	ATOM	316	O	LYS	40	-41.563	81.817	-47.609	1.00	36.18
	ATOM	317	N	GLU	41	-42.215	81.568	-49.748	1.00	40.16
	ATOM	318	CA	GLU	41	-42.948	80.333	-49.516	1.00	42.80
	ATOM	319	CB	GLU	41	-41.952	79.188	-49.369	1.00	45.31
	ATOM	320	CG	GLU	41	-41.012	79.099	-50.565	1.00	56.07
60	ATOM	321	CD	GLU	41	-39.756	78.278	-50.289	1.00	59.42
	ATOM	322	OE1	GLU	41	-39.031	78.583	-49.310	1.00	63.71
	ATOM	323	OE2	GLU	41	-39.489	77.329	-51.059	1.00	63.71
	ATOM	324	C	GLU	41	-43.859	80.417	-48.302	1.00	38.93
	ATOM	325	O	GLU	41	-43.789	79.587	-47.400	1.00	39.03
65	ATOM	326	N	ASN	42	-44.699	81.443	-48.287	1.00	34.69
	ATOM	327	CA	ASN	42	-45.661	81.657	-47.220	1.00	32.96

	ATOM	328	CB	ASN	42	-46.531	80.418	-47.046	1.00	32.65
	ATOM	329	CG	ASN	42	-47.936	80.763	-46.662	1.00	35.50
	ATOM	330	OD1	ASN	42	-48.520	80.140	-45.785	1.00	40.26
	ATOM	331	ND2	ASN	42	-48.497	81.762	-47.325	1.00	29.28
5	ATOM	332	C	ASN	42	-45.066	82.031	-45.871	1.00	31.44
	ATOM	333	O	ASN	42	-45.731	81.924	-44.845	1.00	22.78
	ATOM	334	N	LYS	43	-43.820	82.477	-45.877	1.00	33.26
	ATOM	335	CA	LYS	43	-43.148	82.867	-44.649	1.00	33.68
	ATOM	336	CB	LYS	43	-42.123	81.806	-44.248	1.00	33.55
10	ATOM	337	CG	LYS	43	-42.736	80.492	-43.834	1.00	40.49
	ATOM	338	CD	LYS	43	-41.670	79.466	-43.544	1.00	45.39
	ATOM	339	CE	LYS	43	-40.901	79.120	-44.804	1.00	52.57
	ATOM	340	NZ	LYS	43	-39.919	78.033	-44.554	1.00	62.06
	ATOM	341	C	LYS	43	-42.445	84.189	-44.854	1.00	32.05
15	ATOM	342	O	LYS	43	-42.285	84.639	-45.980	1.00	37.32
	ATOM	343	N	ILE	44	-42.037	84.825	-43.766	1.00	31.31
	ATOM	344	CA	ILE	44	-41.333	86.084	-43.892	1.00	29.52
	ATOM	345	CB	ILE	44	-41.805	87.097	-42.848	1.00	26.38
	ATOM	346	CG2	ILE	44	-41.011	88.380	-42.985	1.00	26.09
20	ATOM	347	CG1	ILE	44	-43.291	87.391	-43.059	1.00	28.61
	ATOM	348	CD1	ILE	44	-43.866	88.394	-42.107	1.00	21.51
	ATOM	349	C	ILE	44	-39.845	85.826	-43.732	1.00	31.60
	ATOM	350	O	ILE	44	-39.401	85.286	-42.722	1.00	34.50
	ATOM	351	N	LEU	45	-39.082	86.197	-44.751	1.00	31.44
25	ATOM	352	CA	LEU	45	-37.639	86.010	-44.745	1.00	31.40
	ATOM	353	CB	LEU	45	-37.176	85.551	-46.125	1.00	33.60
	ATOM	354	CG	LEU	45	-35.667	85.421	-46.316	1.00	34.36
	ATOM	355	CD1	LEU	45	-35.124	84.298	-45.456	1.00	29.00
	ATOM	356	CD2	LEU	45	-35.380	85.161	-47.783	1.00	32.42
30	ATOM	357	C	LEU	45	-36.892	87.282	-44.353	1.00	27.80
	ATOM	358	O	LEU	45	-37.129	88.352	-44.905	1.00	27.94
	ATOM	359	N	VAL	46	-35.987	87.150	-43.392	1.00	28.00
	ATOM	360	CA	VAL	46	-35.187	88.271	-42.922	1.00	31.53
	ATOM	361	CB	VAL	46	-34.777	88.049	-41.457	1.00	27.89
35	ATOM	362	CG1	VAL	46	-33.989	89.230	-40.944	1.00	26.10
	ATOM	363	CG2	VAL	46	-36.006	87.830	-40.618	1.00	23.39
	ATOM	364	C	VAL	46	-33.927	88.403	-43.793	1.00	33.94
	ATOM	365	O	VAL	46	-33.161	87.447	-43.932	1.00	39.68
	ATOM	366	N	LYS	47	-33.714	89.582	-44.373	1.00	33.59
40	ATOM	367	CA	LYS	47	-32.551	89.805	-45.228	1.00	37.05
	ATOM	368	CB	LYS	47	-32.983	90.440	-46.554	1.00	34.49
	ATOM	369	CG	LYS	47	-34.153	89.734	-47.217	1.00	37.53
	ATOM	370	CD	LYS	47	-33.852	88.263	-47.470	1.00	41.69
	ATOM	371	CE	LYS	47	-33.328	88.040	-48.877	1.00	42.95
45	ATOM	372	NZ	LYS	47	-32.273	89.018	-49.226	1.00	44.50
	ATOM	373	C	LYS	47	-31.470	90.667	-44.574	1.00	37.34
	ATOM	374	O	LYS	47	-30.357	90.762	-45.080	1.00	43.04
	ATOM	375	N	GLU	48	-31.806	91.293	-43.455	1.00	36.99
	ATOM	376	CA	GLU	48	-30.868	92.130	-42.714	1.00	37.31
50	ATOM	377	CB	GLU	48	-31.171	93.605	-42.920	1.00	36.53
	ATOM	378	CG	GLU	48	-30.897	94.132	-44.297	1.00	46.74
	ATOM	379	CD	GLU	48	-31.275	95.597	-44.415	1.00	52.39
	ATOM	380	OE1	GLU	48	-31.009	96.358	-43.444	1.00	51.33
	ATOM	381	OE2	GLU	48	-31.832	95.987	-45.474	1.00	54.55
55	ATOM	382	C	GLU	48	-31.062	91.820	-41.250	1.00	36.94
	ATOM	383	O	GLU	48	-32.186	91.864	-40.767	1.00	40.21
	ATOM	384	N	THR	49	-29.992	91.524	-40.524	1.00	36.34
	ATOM	385	CA	THR	49	-30.183	91.208	-39.118	1.00	35.48
	ATOM	386	CB	THR	49	-28.985	90.410	-38.532	1.00	31.16
60	ATOM	387	OG1	THR	49	-28.097	91.300	-37.859	1.00	32.48
	ATOM	388	CG2	THR	49	-28.231	89.694	-39.626	1.00	30.36
	ATOM	389	C	THR	49	-30.422	92.485	-38.307	1.00	34.59
	ATOM	390	O	THR	49	-29.951	93.563	-38.666	1.00	35.13
	ATOM	391	N	GLY	50	-31.181	92.350	-37.223	1.00	32.49
65	ATOM	392	CA	GLY	50	-31.479	93.486	-36.378	1.00	28.17
	ATOM	393	C	GLY	50	-32.649	93.190	-35.468	1.00	29.99

	ATOM	394	O	GLY	50	-32.992	92.038	-35.258	1.00	29.76
	ATOM	395	N	TYR	51	-33.255	94.237	-34.919	1.00	31.68
	ATOM	396	CA	TYR	51	-34.408	94.091	-34.037	1.00	32.88
	ATOM	397	CB	TYR	51	-34.281	95.017	-32.822	1.00	35.97
5	ATOM	398	CG	TYR	51	-33.209	94.582	-31.867	1.00	43.03
	ATOM	399	CD1	TYR	51	-31.867	94.892	-32.103	1.00	44.15
	ATOM	400	CE1	TYR	51	-30.857	94.411	-31.268	1.00	47.81
	ATOM	401	CD2	TYR	51	-33.523	93.786	-30.768	1.00	47.08
	ATOM	402	CE2	TYR	51	-32.526	93.294	-29.926	1.00	51.79
10	ATOM	403	CZ	TYR	51	-31.192	93.606	-30.182	1.00	51.15
	ATOM	404	OH	TYR	51	-30.203	93.075	-29.376	1.00	52.20
	ATOM	405	C	TYR	51	-35.700	94.395	-34.783	1.00	31.97
	ATOM	406	O	TYR	51	-35.845	95.449	-35.397	1.00	30.01
	ATOM	407	N	PHE	52	-36.643	93.461	-34.727	1.00	31.50
15	ATOM	408	CA	PHE	52	-37.910	93.640	-35.415	1.00	30.41
	ATOM	409	CB	PHE	52	-38.081	92.584	-36.509	1.00	31.30
	ATOM	410	CG	PHE	52	-37.009	92.600	-37.554	1.00	29.70
	ATOM	411	CD1	PHE	52	-35.739	92.126	-37.274	1.00	26.20
	ATOM	412	CD2	PHE	52	-37.278	93.081	-38.827	1.00	28.38
20	ATOM	413	CE1	PHE	52	-34.757	92.133	-38.243	1.00	27.17
	ATOM	414	CE2	PHE	52	-36.300	93.091	-39.803	1.00	27.76
	ATOM	415	CZ	PHE	52	-35.038	92.617	-39.512	1.00	27.27
	ATOM	416	C	PHE	52	-39.114	93.556	-34.500	1.00	27.65
	ATOM	417	O	PHE	52	-39.115	92.823	-33.519	1.00	24.85
25	ATOM	418	N	PHE	53	-40.137	94.326	-34.839	1.00	27.18
	ATOM	419	CA	PHE	53	-41.398	94.310	-34.114	1.00	27.05
	ATOM	420	CB	PHE	53	-42.034	95.694	-34.096	1.00	28.76
	ATOM	421	CG	PHE	53	-43.420	95.710	-33.536	1.00	28.00
	ATOM	422	CD1	PHE	53	-43.642	95.511	-32.182	1.00	26.49
30	ATOM	423	CD2	PHE	53	-44.513	95.908	-34.368	1.00	28.04
	ATOM	424	CE1	PHE	53	-44.932	95.509	-31.665	1.00	26.54
	ATOM	425	CE2	PHE	53	-45.806	95.906	-33.854	1.00	25.78
	ATOM	426	CZ	PHE	53	-46.012	95.706	-32.501	1.00	24.39
	ATOM	427	C	PHE	53	-42.245	93.369	-34.966	1.00	27.56
35	ATOM	428	O	PHE	53	-42.453	93.626	-36.148	1.00	25.39
	ATOM	429	N	ILE	54	-42.715	92.278	-34.371	1.00	28.87
	ATOM	430	CA	ILE	54	-43.504	91.287	-35.096	1.00	28.95
	ATOM	431	CB	ILE	54	-42.819	89.908	-35.003	1.00	29.60
	ATOM	432	CG2	ILE	54	-43.470	88.938	-35.976	1.00	27.42
40	ATOM	433	CG1	ILE	54	-41.331	90.055	-35.329	1.00	26.41
	ATOM	434	CD1	ILE	54	-40.491	88.871	-34.961	1.00	28.70
	ATOM	435	C	ILE	54	-44.928	91.197	-34.550	1.00	28.57
	ATOM	436	O	ILE	54	-45.142	91.196	-33.342	1.00	29.17
	ATOM	437	N	TYR	55	-45.908	91.114	-35.440	1.00	26.97
45	ATOM	438	CA	TYR	55	-47.297	91.055	-35.001	1.00	24.22
	ATOM	439	CB	TYR	55	-47.903	92.459	-35.034	1.00	21.82
	ATOM	440	CG	TYR	55	-47.869	93.117	-36.391	1.00	27.68
	ATOM	441	CD1	TYR	55	-48.924	92.971	-37.287	1.00	27.01
	ATOM	442	CE1	TYR	55	-48.882	93.558	-38.539	1.00	27.44
50	ATOM	443	CD2	TYR	55	-46.771	93.872	-36.789	1.00	27.16
	ATOM	444	CE2	TYR	55	-46.720	94.460	-38.043	1.00	25.39
	ATOM	445	CZ	TYR	55	-47.775	94.299	-38.911	1.00	29.02
	ATOM	446	OH	TYR	55	-47.718	94.872	-40.155	1.00	31.82
	ATOM	447	C	TYR	55	-48.143	90.105	-35.821	1.00	25.16
55	ATOM	448	O	TYR	55	-47.763	89.711	-36.901	1.00	28.47
	ATOM	449	N	GLY	56	-49.295	89.727	-35.294	1.00	27.44
	ATOM	450	CA	GLY	56	-50.160	88.819	-36.015	1.00	22.07
	ATOM	451	C	GLY	56	-51.530	88.710	-35.382	1.00	25.83
	ATOM	452	O	GLY	56	-51.667	88.802	-34.166	1.00	26.00
60	ATOM	453	N	GLN	57	-52.548	88.537	-36.215	1.00	22.56
	ATOM	454	CA	GLN	57	-53.917	88.388	-35.748	1.00	20.91
	ATOM	455	CB	GLN	57	-54.683	89.710	-35.803	1.00	17.87
	ATOM	456	CG	GLN	57	-56.156	89.549	-35.445	1.00	16.86
	ATOM	457	CD	GLN	57	-56.922	90.855	-35.409	1.00	22.06
65	ATOM	458	OE1	GLN	57	-56.649	91.723	-34.586	1.00	27.26
	ATOM	459	NE2	GLN	57	-57.893	90.997	-36.302	1.00	20.20

	ATOM	460	C	GLN	57	-54.626	87.370	-36.617	1.00	24.23
	ATOM	461	O	GLN	57	-54.373	87.292	-37.818	1.00	25.16
	ATOM	462	N	VAL	58	-55.509	86.594	-35.995	1.00	24.62
	ATOM	463	CA	VAL	58	-56.287	85.567	-36.677	1.00	26.68
5	ATOM	464	CB	VAL	58	-55.695	84.156	-36.435	1.00	25.55
	ATOM	465	CG1	VAL	58	-56.630	83.101	-36.972	1.00	15.69
	ATOM	466	CG2	VAL	58	-54.339	84.031	-37.095	1.00	25.77
	ATOM	467	C	VAL	58	-57.685	85.587	-36.096	1.00	31.06
	ATOM	468	O	VAL	58	-57.837	85.759	-34.894	1.00	27.74
10	ATOM	469	N	LEU	59	-58.703	85.420	-36.941	1.00	33.82
	ATOM	470	CA	LEU	59	-60.094	85.375	-36.475	1.00	29.85
	ATOM	471	CB	LEU	59	-61.029	86.102	-37.437	1.00	29.62
	ATOM	472	CG	LEU	59	-62.359	86.619	-36.872	1.00	28.11
	ATOM	473	CD1	LEU	59	-63.279	86.980	-38.020	1.00	26.67
15	ATOM	474	CD2	LEU	59	-63.019	85.597	-35.995	1.00	22.45
	ATOM	475	C	LEU	59	-60.513	83.910	-36.398	1.00	30.98
	ATOM	476	O	LEU	59	-60.592	83.224	-37.415	1.00	32.52
	ATOM	477	N	TYR	60	-60.777	83.431	-35.187	1.00	33.01
	ATOM	478	CA	TYR	60	-61.183	82.046	-34.997	1.00	34.51
20	ATOM	479	CB	TYR	60	-60.621	81.511	-33.684	1.00	40.10
	ATOM	480	CG	TYR	60	-59.124	81.511	-33.659	1.00	45.37
	ATOM	481	CD1	TYR	60	-58.416	82.450	-32.909	1.00	47.13
	ATOM	482	CE1	TYR	60	-57.030	82.500	-32.958	1.00	50.28
	ATOM	483	CD2	TYR	60	-58.409	80.619	-34.449	1.00	44.35
25	ATOM	484	CE2	TYR	60	-57.030	80.657	-34.508	1.00	48.81
	ATOM	485	CZ	TYR	60	-56.347	81.599	-33.767	1.00	51.91
	ATOM	486	OH	TYR	60	-54.979	81.655	-33.868	1.00	53.77
	ATOM	487	C	TYR	60	-62.690	81.854	-35.018	1.00	35.49
	ATOM	488	O	TYR	60	-63.421	82.458	-34.240	1.00	36.13
30	ATOM	489	N	THR	61	-63.146	81.004	-35.926	1.00	36.19
	ATOM	490	CA	THR	61	-64.558	80.702	-36.056	1.00	35.36
	ATOM	491	CB	THR	61	-65.075	81.100	-37.440	1.00	34.22
	ATOM	492	OG1	THR	61	-64.249	80.505	-38.445	1.00	34.22
	ATOM	493	CG2	THR	61	-65.060	82.608	-37.598	1.00	25.45
35	ATOM	494	C	THR	61	-64.706	79.200	-35.855	1.00	37.82
	ATOM	495	O	THR	61	-65.566	78.553	-36.445	1.00	43.70
	ATOM	496	N	ASP	62	-63.839	78.664	-35.007	1.00	37.64
	ATOM	497	CA	ASP	62	-63.795	77.251	-34.676	1.00	38.08
	ATOM	498	CB	ASP	62	-62.346	76.788	-34.794	1.00	38.42
40	ATOM	499	CG	ASP	62	-62.191	75.301	-34.654	1.00	41.25
	ATOM	500	OD1	ASP	62	-61.358	74.727	-35.395	1.00	38.28
	ATOM	501	OD2	ASP	62	-62.890	74.718	-33.800	1.00	41.37
	ATOM	502	C	ASP	62	-64.300	77.107	-33.243	1.00	40.90
	ATOM	503	O	ASP	62	-63.995	77.946	-32.406	1.00	46.73
45	ATOM	504	N	LYS	63	-65.067	76.063	-32.943	1.00	42.62
	ATOM	505	CA	LYS	63	-65.583	75.913	-31.582	1.00	40.92
	ATOM	506	CB	LYS	63	-67.052	75.488	-31.614	1.00	42.79
	ATOM	507	CG	LYS	63	-67.325	74.186	-32.335	1.00	45.91
	ATOM	508	CD	LYS	63	-68.830	73.973	-32.495	1.00	46.22
50	ATOM	509	CE	LYS	63	-69.461	75.092	-33.345	1.00	49.97
	ATOM	510	NZ	LYS	63	-70.965	75.074	-33.359	1.00	49.98
	ATOM	511	C	LYS	63	-64.797	74.966	-30.682	1.00	42.59
	ATOM	512	O	LYS	63	-65.307	74.508	-29.667	1.00	42.46
	ATOM	513	N	THR	64	-63.544	74.712	-31.037	1.00	44.06
55	ATOM	514	CA	THR	64	-62.682	73.819	-30.276	1.00	44.94
	ATOM	515	CB	THR	64	-61.404	73.490	-31.088	1.00	43.79
	ATOM	516	OG1	THR	64	-61.778	72.963	-32.364	1.00	46.20
	ATOM	517	CG2	THR	64	-60.548	72.452	-30.371	1.00	42.80
	ATOM	518	C	THR	64	-62.272	74.296	-28.873	1.00	45.82
60	ATOM	519	O	THR	64	-61.120	74.175	-28.496	1.00	53.10
	ATOM	520	N	TYR	65	-63.207	74.824	-28.099	1.00	45.11
	ATOM	521	CA	TYR	65	-62.948	75.280	-26.725	1.00	45.72
	ATOM	522	CB	TYR	65	-62.836	74.068	-25.780	1.00	40.46
	ATOM	523	CG	TYR	65	-61.436	73.556	-25.524	1.00	41.90
65	ATOM	524	CD1	TYR	65	-60.649	74.090	-24.501	1.00	42.47
	ATOM	525	CE1	TYR	65	-59.343	73.624	-24.266	1.00	41.83

	ATOM	526	CD2	TYR	65	-60.887	72.539	-26.311	1.00	41.65
	ATOM	527	CE2	TYR	65	-59.582	72.065	-26.089	1.00	40.28
	ATOM	528	CZ	TYR	65	-58.821	72.614	-25.065	1.00	41.90
	ATOM	529	OH	TYR	65	-57.546	72.156	-24.844	1.00	43.84
5	ATOM	530	C	TYR	65	-61.759	76.225	-26.491	1.00	45.31
	ATOM	531	O	TYR	65	-61.792	77.061	-25.579	1.00	49.74
	ATOM	532	N	ALA	66	-60.712	76.096	-27.295	1.00	42.32
	ATOM	533	CA	ALA	66	-59.534	76.942	-27.158	1.00	39.57
	ATOM	534	CB	ALA	66	-58.643	76.424	-26.050	1.00	35.36
10	ATOM	535	C	ALA	66	-58.771	76.974	-28.470	1.00	39.85
	ATOM	536	O	ALA	66	-58.355	75.936	-28.986	1.00	37.60
	ATOM	537	N	MET	67	-58.605	78.172	-29.017	1.00	41.88
	ATOM	538	CA	MET	67	-57.886	78.343	-30.270	1.00	41.48
	ATOM	539	CB	MET	67	-58.823	78.847	-31.375	1.00	39.98
15	ATOM	540	CG	MET	67	-59.872	77.843	-31.833	1.00	37.31
	ATOM	541	SD	MET	67	-59.160	76.322	-32.476	1.00	35.46
	ATOM	542	CE	MET	67	-58.721	76.803	-34.148	1.00	34.06
	ATOM	543	C	MET	67	-56.752	79.332	-30.064	1.00	39.50
	ATOM	544	O	MET	67	-56.703	80.037	-29.054	1.00	38.44
20	ATOM	545	N	GLY	68	-55.840	79.377	-31.027	1.00	39.84
	ATOM	546	CA	GLY	68	-54.717	80.288	-30.931	1.00	37.61
	ATOM	547	C	GLY	68	-53.678	80.022	-31.994	1.00	33.76
	ATOM	548	O	GLY	68	-53.772	79.044	-32.726	1.00	35.58
	ATOM	549	N	HIS	69	-52.691	80.902	-32.091	1.00	35.07
25	ATOM	550	CA	HIS	69	-51.631	80.738	-33.072	1.00	33.70
	ATOM	551	CB	HIS	69	-51.864	81.649	-34.292	1.00	37.02
	ATOM	552	CG	HIS	69	-52.066	83.095	-33.957	1.00	34.65
	ATOM	553	CD2	HIS	69	-51.248	84.163	-34.098	1.00	35.63
	ATOM	554	ND1	HIS	69	-53.238	83.582	-33.422	1.00	37.68
30	ATOM	555	CE1	HIS	69	-53.136	84.887	-33.248	1.00	32.64
	ATOM	556	NE2	HIS	69	-51.937	85.265	-33.651	1.00	35.90
	ATOM	557	C	HIS	69	-50.262	81.013	-32.468	1.00	33.98
	ATOM	558	O	HIS	69	-50.155	81.506	-31.356	1.00	30.94
	ATOM	559	N	LEU	70	-49.215	80.682	-33.210	1.00	35.35
35	ATOM	560	CA	LEU	70	-47.849	80.892	-32.752	1.00	30.76
	ATOM	561	CB	LEU	70	-47.144	79.554	-32.560	1.00	29.59
	ATOM	562	CG	LEU	70	-47.916	78.389	-31.948	1.00	30.71
	ATOM	563	CD1	LEU	70	-47.105	77.125	-32.099	1.00	25.60
	ATOM	564	CD2	LEU	70	-48.219	78.674	-30.497	1.00	27.39
40	ATOM	565	C	LEU	70	-47.059	81.676	-33.795	1.00	33.19
	ATOM	566	O	LEU	70	-47.187	81.428	-34.994	1.00	37.07
	ATOM	567	N	ILE	71	-46.261	82.639	-33.349	1.00	30.93
	ATOM	568	CA	ILE	71	-45.413	83.370	-34.276	1.00	27.94
	ATOM	569	CB	ILE	71	-45.333	84.862	-33.943	1.00	28.22
45	ATOM	570	CG2	ILE	71	-44.192	85.507	-34.702	1.00	24.54
	ATOM	571	CG1	ILE	71	-46.644	85.539	-34.326	1.00	26.34
	ATOM	572	CD1	ILE	71	-46.746	86.964	-33.860	1.00	37.50
	ATOM	573	C	ILE	71	-44.074	82.700	-34.014	1.00	30.32
	ATOM	574	O	ILE	71	-43.545	82.792	-32.912	1.00	23.07
50	ATOM	575	N	GLN	72	-43.542	82.000	-35.012	1.00	29.74
	ATOM	576	CA	GLN	72	-42.288	81.288	-34.818	1.00	27.85
	ATOM	577	CB	GLN	72	-42.503	79.810	-35.107	1.00	26.70
	ATOM	578	CG	GLN	72	-43.691	79.249	-34.380	1.00	28.28
	ATOM	579	CD	GLN	72	-43.787	77.756	-34.506	1.00	33.11
55	ATOM	580	OE1	GLN	72	-43.781	77.219	-35.610	1.00	41.22
	ATOM	581	NE2	GLN	72	-43.880	77.069	-33.375	1.00	30.18
	ATOM	582	C	GLN	72	-41.110	81.792	-35.625	1.00	29.85
	ATOM	583	O	GLN	72	-41.275	82.440	-36.661	1.00	28.20
	ATOM	584	N	ARG	73	-39.918	81.467	-35.139	1.00	30.90
60	ATOM	585	CA	ARG	73	-38.679	81.863	-35.789	1.00	30.79
	ATOM	586	CB	ARG	73	-37.908	82.805	-34.875	1.00	29.18
	ATOM	587	CG	ARG	73	-36.567	83.208	-35.409	1.00	28.66
	ATOM	588	CD	ARG	73	-35.671	83.635	-34.284	1.00	29.25
	ATOM	589	NE	ARG	73	-34.375	84.083	-34.771	1.00	35.78
65	ATOM	590	CZ	ARG	73	-33.291	84.181	-34.016	1.00	35.27
	ATOM	591	NH1	ARG	73	-33.343	83.855	-32.732	1.00	34.95

	ATOM	592	NH2	ARG	73	-32.160	84.609	-34.546	1.00	33.33
	ATOM	593	C	ARG	73	-37.800	80.658	-36.119	1.00	32.42
	ATOM	594	O	ARG	73	-37.532	79.831	-35.255	1.00	34.19
	ATOM	595	N	LYS	74	-37.374	80.557	-37.374	1.00	35.90
5	ATOM	596	CA	LYS	74	-36.483	79.479	-37.801	1.00	38.57
	ATOM	597	CB	LYS	74	-36.876	78.942	-39.180	1.00	43.52
	ATOM	598	CG	LYS	74	-38.139	78.097	-39.199	1.00	52.89
	ATOM	599	CD	LYS	74	-38.506	77.626	-40.626	1.00	58.14
	ATOM	600	CE	LYS	74	-39.788	76.792	-40.613	1.00	56.75
10	ATOM	601	NZ	LYS	74	-40.194	76.358	-41.983	1.00	63.98
	ATOM	602	C	LYS	74	-35.081	80.068	-37.895	1.00	40.11
	ATOM	603	O	LYS	74	-34.781	80.825	-38.827	1.00	37.55
	ATOM	604	N	LYS	75	-34.228	79.725	-36.936	1.00	34.48
	ATOM	605	CA	LYS	75	-32.861	80.232	-36.908	1.00	34.24
15	ATOM	606	CB	LYS	75	-32.185	79.839	-35.592	1.00	36.16
	ATOM	607	CG	LYS	75	-32.875	80.292	-34.308	1.00	35.99
	ATOM	608	CD	LYS	75	-32.074	79.800	-33.095	1.00	40.91
	ATOM	609	CE	LYS	75	-32.697	80.242	-31.778	1.00	46.16
	ATOM	610	NZ	LYS	75	-31.985	79.741	-30.550	1.00	46.03
20	ATOM	611	C	LYS	75	-32.031	79.679	-38.071	1.00	30.92
	ATOM	612	O	LYS	75	-32.148	78.515	-38.407	1.00	27.47
	ATOM	613	N	VAL	76	-31.192	80.518	-38.673	1.00	32.82
	ATOM	614	CA	VAL	76	-30.321	80.091	-39.772	1.00	31.51
	ATOM	615	CB	VAL	76	-29.723	81.270	-40.561	1.00	29.19
25	ATOM	616	CG1	VAL	76	-29.595	80.897	-42.010	1.00	27.09
	ATOM	617	CG2	VAL	76	-30.547	82.495	-40.384	1.00	34.94
	ATOM	618	C	VAL	76	-29.132	79.378	-39.160	1.00	33.40
	ATOM	619	O	VAL	76	-28.661	78.371	-39.677	1.00	32.92
	ATOM	620	N	HIS	77	-28.643	79.938	-38.058	1.00	34.88
30	ATOM	621	CA	HIS	77	-27.503	79.396	-37.342	1.00	34.59
	ATOM	622	CB	HIS	77	-26.501	80.511	-37.028	1.00	35.92
	ATOM	623	CG	HIS	77	-26.002	81.237	-38.239	1.00	34.17
	ATOM	624	CD2	HIS	77	-25.509	82.487	-38.388	1.00	31.44
	ATOM	625	ND1	HIS	77	-25.947	80.654	-39.489	1.00	35.05
35	ATOM	626	CE1	HIS	77	-25.444	81.511	-40.356	1.00	29.84
	ATOM	627	NE2	HIS	77	-25.168	82.630	-39.713	1.00	38.35
	ATOM	628	C	HIS	77	-27.986	78.736	-36.061	1.00	35.63
	ATOM	629	O	HIS	77	-28.862	79.269	-35.379	1.00	40.24
	ATOM	630	N	VAL	78	-27.399	77.590	-35.720	1.00	37.37
40	ATOM	631	CA	VAL	78	-27.840	76.866	-34.542	1.00	41.32
	ATOM	632	CB	VAL	78	-28.419	75.500	-34.958	1.00	43.10
	ATOM	633	CG1	VAL	78	-29.011	74.800	-33.759	1.00	45.01
	ATOM	634	CG2	VAL	78	-29.508	75.699	-35.998	1.00	44.98
	ATOM	635	C	VAL	78	-26.897	76.681	-33.346	1.00	43.02
45	ATOM	636	O	VAL	78	-27.167	77.232	-32.280	1.00	49.79
	ATOM	637	N	PHE	79	-25.817	75.922	-33.474	1.00	41.33
	ATOM	638	CA	PHE	79	-24.911	75.722	-32.316	1.00	45.00
	ATOM	639	CB	PHE	79	-24.664	77.017	-31.516	1.00	40.51
	ATOM	640	CG	PHE	79	-24.151	78.158	-32.328	1.00	39.52
50	ATOM	641	CD1	PHE	79	-24.992	79.214	-32.664	1.00	36.80
	ATOM	642	CD2	PHE	79	-22.830	78.185	-32.753	1.00	37.89
	ATOM	643	CE1	PHE	79	-24.524	80.281	-33.411	1.00	37.52
	ATOM	644	CE2	PHE	79	-22.354	79.250	-33.501	1.00	35.59
	ATOM	645	CZ	PHE	79	-23.201	80.301	-33.832	1.00	37.58
55	ATOM	646	C	PHE	79	-25.407	74.704	-31.289	1.00	43.50
	ATOM	647	O	PHE	79	-26.503	74.834	-30.742	1.00	37.85
	ATOM	648	N	GLY	80	-24.566	73.715	-31.009	1.00	46.65
	ATOM	649	CA	GLY	80	-24.884	72.698	-30.025	1.00	48.38
	ATOM	650	C	GLY	80	-26.262	72.075	-30.083	1.00	47.89
60	ATOM	651	O	GLY	80	-26.725	71.657	-31.148	1.00	51.15
	ATOM	652	N	ASP	81	-26.918	72.015	-28.927	1.00	45.07
	ATOM	653	CA	ASP	81	-28.235	71.412	-28.829	1.00	43.99
	ATOM	654	CB	ASP	81	-28.340	70.587	-27.540	1.00	45.66
	ATOM	655	CG	ASP	81	-28.350	71.446	-26.285	1.00	47.49
65	ATOM	656	OD1	ASP	81	-28.014	72.649	-26.362	1.00	51.90
	ATOM	657	OD2	ASP	81	-28.686	70.913	-25.210	1.00	48.08

	ATOM	658	C	ASP	81	-29.382	72.403	-28.907	1.00	42.29
	ATOM	659	O	ASP	81	-30.479	72.121	-28.426	1.00	39.94
	ATOM	660	N	GLU	82	-29.134	73.562	-29.509	1.00	39.91
	ATOM	661	CA	GLU	82	-30.184	74.567	-29.665	1.00	36.58
5	ATOM	662	CB	GLU	82	-29.657	75.839	-30.332	1.00	34.91
	ATOM	663	CG	GLU	82	-28.896	76.801	-29.477	1.00	37.82
	ATOM	664	CD	GLU	82	-28.959	78.211	-30.043	1.00	40.75
	ATOM	665	OE1	GLU	82	-28.811	78.372	-31.270	1.00	38.15
	ATOM	666	OE2	GLU	82	-29.162	79.164	-29.263	1.00	47.99
10	ATOM	667	C	GLU	82	-31.266	74.022	-30.587	1.00	36.58
	ATOM	668	O	GLU	82	-30.985	73.235	-31.486	1.00	38.40
	ATOM	669	N	LEU	83	-32.502	74.442	-30.359	1.00	33.83
	ATOM	670	CA	LEU	83	-33.606	74.056	-31.222	1.00	32.70
	ATOM	671	CB	LEU	83	-34.930	74.085	-30.458	1.00	30.78
15	ATOM	672	CG	LEU	83	-35.411	72.789	-29.809	1.00	29.26
	ATOM	673	CD1	LEU	83	-34.273	72.074	-29.125	1.00	29.60
	ATOM	674	CD2	LEU	83	-36.507	73.112	-28.826	1.00	28.65
	ATOM	675	C	LEU	83	-33.584	75.162	-32.255	1.00	33.88
	ATOM	676	O	LEU	83	-33.489	76.330	-31.896	1.00	38.46
20	ATOM	677	N	SER	84	-33.654	74.812	-33.531	1.00	34.94
	ATOM	678	CA	SER	84	-33.613	75.833	-34.567	1.00	39.77
	ATOM	679	CB	SER	84	-33.122	75.236	-35.884	1.00	38.92
	ATOM	680	OG	SER	84	-33.962	74.185	-36.293	1.00	43.11
	ATOM	681	C	SER	84	-34.952	76.532	-34.779	1.00	39.89
25	ATOM	682	O	SER	84	-35.034	77.551	-35.465	1.00	42.52
	ATOM	683	N	LEU	85	-36.001	75.985	-34.182	1.00	40.92
	ATOM	684	CA	LEU	85	-37.325	76.566	-34.297	1.00	34.66
	ATOM	685	CB	LEU	85	-38.324	75.501	-34.751	1.00	35.12
	ATOM	686	CG	LEU	85	-39.738	75.874	-35.218	1.00	36.92
30	ATOM	687	CD1	LEU	85	-40.553	76.393	-34.059	1.00	38.24
	ATOM	688	CD2	LEU	85	-39.657	76.906	-36.333	1.00	36.08
	ATOM	689	C	LEU	85	-37.699	77.099	-32.926	1.00	35.30
	ATOM	690	O	LEU	85	-37.831	76.346	-31.965	1.00	32.85
	ATOM	691	N	VAL	86	-37.843	78.412	-32.836	1.00	33.26
35	ATOM	692	CA	VAL	86	-38.204	79.052	-31.581	1.00	34.57
	ATOM	693	CB	VAL	86	-37.190	80.163	-31.184	1.00	35.39
	ATOM	694	CG1	VAL	86	-37.688	80.921	-29.969	1.00	31.44
	ATOM	695	CG2	VAL	86	-35.834	79.551	-30.890	1.00	38.08
	ATOM	696	C	VAL	86	-39.560	79.700	-31.731	1.00	32.85
40	ATOM	697	O	VAL	86	-39.821	80.379	-32.719	1.00	35.10
	ATOM	698	N	THR	87	-40.436	79.487	-30.765	1.00	33.41
	ATOM	699	CA	THR	87	-41.725	80.132	-30.846	1.00	33.56
	ATOM	700	CB	THR	87	-42.889	79.134	-30.557	1.00	32.70
	ATOM	701	OG1	THR	87	-43.894	79.767	-29.755	1.00	34.29
45	ATOM	702	CG2	THR	87	-42.372	77.887	-29.890	1.00	34.95
	ATOM	703	C	THR	87	-41.762	81.351	-29.917	1.00	30.57
	ATOM	704	O	THR	87	-41.632	81.235	-28.701	1.00	28.08
	ATOM	705	N	LEU	88	-41.872	82.525	-30.533	1.00	28.42
	ATOM	706	CA	LEU	88	-41.967	83.786	-29.822	1.00	31.31
50	ATOM	707	CB	LEU	88	-41.371	84.933	-30.649	1.00	28.21
	ATOM	708	CG	LEU	88	-39.993	85.049	-31.308	1.00	29.12
	ATOM	709	CD1	LEU	88	-39.098	83.931	-30.904	1.00	31.12
	ATOM	710	CD2	LEU	88	-40.166	85.085	-32.808	1.00	28.72
	ATOM	711	C	LEU	88	-43.472	84.069	-29.678	1.00	41.15
55	ATOM	712	O	LEU	88	-44.276	83.688	-30.532	1.00	52.30
	ATOM	713	N	PHE	89	-43.880	84.712	-28.600	1.00	39.23
	ATOM	714	CA	PHE	89	-45.294	85.075	-28.479	1.00	42.27
	ATOM	715	CB	PHE	89	-45.610	86.065	-29.603	1.00	35.63
	ATOM	716	CG	PHE	89	-44.508	87.072	-29.809	1.00	35.06
60	ATOM	717	CD1	PHE	89	-44.081	87.408	-31.089	1.00	31.56
	ATOM	718	CD2	PHE	89	-43.811	87.590	-28.704	1.00	30.96
	ATOM	719	CE1	PHE	89	-42.964	88.232	-31.270	1.00	33.76
	ATOM	720	CE2	PHE	89	-42.699	88.408	-28.872	1.00	27.38
	ATOM	721	CZ	PHE	89	-42.272	88.728	-30.155	1.00	30.29
65	ATOM	722	C	PHE	89	-46.356	83.956	-28.382	1.00	39.99
	ATOM	723	O	PHE	89	-46.350	83.186	-27.417	1.00	45.34

	ATOM	724	N	ARG	90	-47.278	83.856	-29.329	1.00	38.92
	ATOM	725	CA	ARG	90	-48.329	82.830	-29.203	1.00	41.66
	ATOM	726	CB	ARG	90	-47.741	81.493	-28.704	1.00	39.24
	ATOM	727	CG	ARG	90	-48.648	80.703	-27.739	1.00	29.41
5	ATOM	728	CD	ARG	90	-48.016	79.368	-27.301	1.00	31.13
	ATOM	729	NE	ARG	90	-48.737	78.752	-26.185	1.00	31.75
	ATOM	730	CZ	ARG	90	-48.247	78.586	-24.954	1.00	28.86
	ATOM	731	NH1	ARG	90	-47.020	78.976	-24.650	1.00	25.48
	ATOM	732	NH2	ARG	90	-49.002	78.053	-24.008	1.00	29.84
10	ATOM	733	C	ARG	90	-49.484	83.214	-28.258	1.00	39.23
	ATOM	734	O	ARG	90	-49.257	83.620	-27.120	1.00	31.33
	ATOM	735	N	CYS	91	-50.723	83.084	-28.740	1.00	40.12
	ATOM	736	CA	CYS	91	-51.880	83.373	-27.908	1.00	37.09
	ATOM	737	C	CYS	91	-53.015	82.391	-27.972	1.00	37.21
15	ATOM	738	O	CYS	91	-53.068	81.540	-28.851	1.00	37.46
	ATOM	739	CB	CYS	91	-52.436	84.756	-28.160	1.00	40.13
	ATOM	740	SG	CYS	91	-53.095	85.276	-29.784	1.00	41.64
	ATOM	741	N	ILE	92	-53.930	82.531	-27.019	1.00	34.02
	ATOM	742	CA	ILE	92	-55.074	81.638	-26.905	1.00	33.22
20	ATOM	743	CB	ILE	92	-54.852	80.639	-25.751	1.00	32.88
	ATOM	744	CG2	ILE	92	-55.943	79.590	-25.751	1.00	32.61
	ATOM	745	CG1	ILE	92	-53.486	79.966	-25.907	1.00	34.64
	ATOM	746	CD1	ILE	92	-53.071	79.105	-24.725	1.00	32.30
	ATOM	747	C	ILE	92	-56.381	82.377	-26.650	1.00	30.78
25	ATOM	748	O	ILE	92	-56.386	83.461	-26.103	1.00	33.43
	ATOM	749	N	GLN	93	-57.490	81.787	-27.079	1.00	29.35
	ATOM	750	CA	GLN	93	-58.811	82.357	-26.867	1.00	28.69
	ATOM	751	CB	GLN	93	-59.308	83.120	-28.095	1.00	28.17
	ATOM	752	CG	GLN	93	-58.856	84.575	-28.246	1.00	25.87
30	ATOM	753	CD	GLN	93	-59.292	85.474	-27.110	1.00	24.22
	ATOM	754	OE1	GLN	93	-58.596	85.604	-26.115	1.00	28.90
	ATOM	755	NE2	GLN	93	-60.454	86.097	-27.254	1.00	25.29
	ATOM	756	C	GLN	93	-59.760	81.205	-26.622	1.00	31.44
	ATOM	757	O	GLN	93	-59.812	80.275	-27.412	1.00	32.07
35	ATOM	758	N	ASN	94	-60.489	81.239	-25.515	1.00	35.77
	ATOM	759	CA	ASN	94	-61.458	80.192	-25.256	1.00	33.56
	ATOM	760	CB	ASN	94	-62.002	80.288	-23.829	1.00	34.70
	ATOM	761	CG	ASN	94	-61.085	79.660	-22.811	1.00	33.01
	ATOM	762	OD1	ASN	94	-60.674	78.521	-22.960	1.00	30.56
40	ATOM	763	ND2	ASN	94	-60.774	80.394	-21.761	1.00	36.61
	ATOM	764	C	ASN	94	-62.580	80.430	-26.266	1.00	35.80
	ATOM	765	O	ASN	94	-62.911	81.576	-26.580	1.00	36.63
	ATOM	766	N	MET	95	-63.158	79.356	-26.787	1.00	35.73
	ATOM	767	CA	MET	95	-64.227	79.484	-27.769	1.00	34.73
45	ATOM	768	CB	MET	95	-63.886	78.655	-29.011	1.00	32.97
	ATOM	769	CG	MET	95	-62.532	78.960	-29.636	1.00	27.35
	ATOM	770	SD	MET	95	-62.291	80.699	-30.051	1.00	25.14
	ATOM	771	CE	MET	95	-63.337	80.915	-31.450	1.00	15.55
	ATOM	772	C	MET	95	-65.586	79.046	-27.219	1.00	36.36
50	ATOM	773	O	MET	95	-65.657	78.212	-26.312	1.00	40.70
	ATOM	774	N	PRO	96	-66.679	79.627	-27.740	1.00	36.06
	ATOM	775	CD	PRO	96	-66.693	80.867	-28.517	1.00	35.32
	ATOM	776	CA	PRO	96	-68.040	79.293	-27.318	1.00	40.00
	ATOM	777	CB	PRO	96	-68.840	80.546	-27.660	1.00	33.97
55	ATOM	778	CG	PRO	96	-67.819	81.592	-27.864	1.00	37.05
	ATOM	779	C	PRO	96	-68.495	78.111	-28.171	1.00	46.34
	ATOM	780	O	PRO	96	-67.849	77.778	-29.164	1.00	47.59
	ATOM	781	N	GLU	97	-69.609	77.485	-27.804	1.00	52.14
	ATOM	782	CA	GLU	97	-70.111	76.347	-28.564	1.00	57.27
60	ATOM	783	CB	GLU	97	-70.949	75.435	-27.668	1.00	62.29
	ATOM	784	CG	GLU	97	-71.056	73.993	-28.154	1.00	72.31
	ATOM	785	CD	GLU	97	-70.085	73.058	-27.418	1.00	79.34
	ATOM	786	OE1	GLU	97	-70.216	72.915	-26.163	1.00	84.94
	ATOM	787	OE2	GLU	97	-69.191	72.472	-28.091	1.00	81.39
65	ATOM	788	C	GLU	97	-70.981	76.856	-29.708	1.00	56.67
	ATOM	789	O	GLU	97	-70.980	76.293	-30.806	1.00	59.18

	ATOM	790	N	THR	98	-71.705	77.939	-29.449	1.00	57.55
	ATOM	791	CA	THR	98	-72.618	78.515	-30.430	1.00	59.71
	ATOM	792	CB	THR	98	-73.506	79.566	-29.777	1.00	59.24
	ATOM	793	OG1	THR	98	-72.663	80.588	-29.208	1.00	68.81
5	ATOM	794	CG2	THR	98	-74.387	78.917	-28.690	1.00	56.15
	ATOM	795	C	THR	98	-71.997	79.145	-31.676	1.00	60.32
	ATOM	796	O	THR	98	-71.712	78.438	-32.650	1.00	65.03
	ATOM	797	N	LEU	99	-71.808	80.465	-31.659	1.00	54.94
	ATOM	798	CA	LEU	99	-71.250	81.175	-32.813	1.00	51.13
10	ATOM	799	CB	LEU	99	-72.134	82.373	-33.152	1.00	50.56
	ATOM	800	CG	LEU	99	-73.480	82.065	-33.802	1.00	52.80
	ATOM	801	CD1	LEU	99	-74.412	83.270	-33.686	1.00	48.38
	ATOM	802	CD2	LEU	99	-73.253	81.687	-35.258	1.00	50.80
	ATOM	803	C	LEU	99	-69.805	81.645	-32.619	1.00	50.20
15	ATOM	804	O	LEU	99	-69.552	82.834	-32.377	1.00	50.50
	ATOM	805	N	PRO	100	-68.834	80.725	-32.761	1.00	47.97
	ATOM	806	CD	PRO	100	-68.984	79.341	-33.239	1.00	45.68
	ATOM	807	CA	PRO	100	-67.417	81.055	-32.590	1.00	46.54
	ATOM	808	CB	PRO	100	-66.704	79.764	-32.997	1.00	45.50
20	ATOM	809	CG	PRO	100	-67.723	78.705	-32.729	1.00	45.80
	ATOM	810	C	PRO	100	-66.976	82.234	-33.440	1.00	46.78
	ATOM	811	O	PRO	100	-67.119	82.220	-34.657	1.00	52.13
	ATOM	812	N	ASN	101	-66.440	83.254	-32.793	1.00	42.71
	ATOM	813	CA	ASN	101	-65.948	84.432	-33.481	1.00	41.37
25	ATOM	814	CB	ASN	101	-67.110	85.299	-33.951	1.00	43.24
	ATOM	815	CG	ASN	101	-67.679	84.843	-35.280	1.00	45.93
	ATOM	816	OD1	ASN	101	-67.036	84.974	-36.323	1.00	45.33
	ATOM	817	ND2	ASN	101	-68.893	84.297	-35.249	1.00	48.11
	ATOM	818	C	ASN	101	-65.074	85.209	-32.517	1.00	42.66
30	ATOM	819	O	ASN	101	-65.526	86.167	-31.887	1.00	44.79
	ATOM	820	N	ASN	102	-63.816	84.800	-32.386	1.00	40.46
	ATOM	821	CA	ASN	102	-62.949	85.507	-31.473	1.00	37.90
	ATOM	822	CB	ASN	102	-62.425	84.567	-30.398	1.00	34.01
	ATOM	823	CG	ASN	102	-63.374	84.466	-29.220	1.00	32.44
35	ATOM	824	OD1	ASN	102	-63.949	85.463	-28.793	1.00	29.07
	ATOM	825	ND2	ASN	102	-63.536	83.265	-28.686	1.00	33.79
	ATOM	826	C	ASN	102	-61.812	86.342	-32.030	1.00	39.20
	ATOM	827	O	ASN	102	-61.818	87.556	-31.868	1.00	46.07
	ATOM	828	N	SER	103	-60.838	85.743	-32.687	1.00	35.42
40	ATOM	829	CA	SER	103	-59.720	86.562	-33.187	1.00	40.28
	ATOM	830	CB	SER	103	-60.215	87.754	-34.042	1.00	40.68
	ATOM	831	OG	SER	103	-60.193	88.989	-33.335	1.00	31.76
	ATOM	832	C	SER	103	-58.859	87.084	-32.012	1.00	35.24
	ATOM	833	O	SER	103	-59.338	87.755	-31.100	1.00	24.65
45	ATOM	834	N	CYS	104	-57.581	86.728	-32.043	1.00	32.88
	ATOM	835	CA	CYS	104	-56.650	87.127	-31.019	1.00	31.77
	ATOM	836	C	CYS	104	-55.455	87.814	-31.703	1.00	30.58
	ATOM	837	O	CYS	104	-54.955	87.346	-32.722	1.00	29.23
	ATOM	838	CB	CYS	104	-56.190	85.896	-30.190	1.00	28.43
50	ATOM	839	SG	CYS	104	-54.764	86.387	-29.208	1.00	46.07
	ATOM	840	N	TYR	105	-55.026	88.949	-31.154	1.00	30.34
	ATOM	841	CA	TYR	105	-53.876	89.690	-31.671	1.00	26.12
	ATOM	842	CB	TYR	105	-54.226	91.165	-31.858	1.00	27.03
	ATOM	843	CG	TYR	105	-53.049	92.062	-32.208	1.00	25.61
55	ATOM	844	CD1	TYR	105	-52.780	92.425	-33.531	1.00	24.90
	ATOM	845	CE1	TYR	105	-51.705	93.242	-33.851	1.00	26.08
	ATOM	846	CD2	TYR	105	-52.200	92.545	-31.217	1.00	25.41
	ATOM	847	CE2	TYR	105	-51.120	93.360	-31.531	1.00	28.91
	ATOM	848	CZ	TYR	105	-50.879	93.705	-32.848	1.00	27.58
60	ATOM	849	OH	TYR	105	-49.815	94.525	-33.153	1.00	29.89
	ATOM	850	C	TYR	105	-52.741	89.575	-30.657	1.00	29.20
	ATOM	851	O	TYR	105	-52.972	89.563	-29.452	1.00	33.10
	ATOM	852	N	SER	106	-51.512	89.490	-31.139	1.00	30.48
	ATOM	853	CA	SER	106	-50.374	89.391	-30.239	1.00	29.22
65	ATOM	854	CB	SER	106	-50.181	87.939	-29.795	1.00	28.30
	ATOM	855	OG	SER	106	-49.113	87.810	-28.877	1.00	28.62

	ATOM	856	C	SER	106	-49.145	89.890	-30.979	1.00	29.94
	ATOM	857	O	SER	106	-49.016	89.661	-32.175	1.00	29.72
	ATOM	858	N	ALA	107	-48.259	90.588	-30.274	1.00	27.64
	ATOM	859	CA	ALA	107	-47.043	91.112	-30.883	1.00	24.33
5	ATOM	860	CB	ALA	107	-47.327	92.442	-31.557	1.00	21.30
	ATOM	861	C	ALA	107	-45.926	91.277	-29.865	1.00	25.56
	ATOM	862	O	ALA	107	-46.151	91.240	-28.662	1.00	28.03
	ATOM	863	N	GLY	108	-44.713	91.461	-30.360	1.00	26.07
	ATOM	864	CA	GLY	108	-43.576	91.644	-29.481	1.00	25.49
10	ATOM	865	C	GLY	108	-42.330	91.926	-30.288	1.00	29.03
	ATOM	866	O	GLY	108	-42.371	91.902	-31.511	1.00	32.20
	ATOM	867	N	ILE	109	-41.222	92.193	-29.612	1.00	29.34
	ATOM	868	CA	ILE	109	-39.965	92.479	-30.288	1.00	27.71
	ATOM	869	CB	ILE	109	-39.298	93.733	-29.699	1.00	27.48
15	ATOM	870	CG2	ILE	109	-37.982	94.005	-30.397	1.00	26.46
	ATOM	871	CG1	ILE	109	-40.238	94.928	-29.851	1.00	26.16
	ATOM	872	CD1	ILE	109	-39.761	96.174	-29.162	1.00	29.07
	ATOM	873	C	ILE	109	-39.023	91.300	-30.138	1.00	28.66
	ATOM	874	O	ILE	109	-38.988	90.653	-29.094	1.00	30.59
20	ATOM	875	N	ALA	110	-38.262	91.014	-31.185	1.00	25.60
	ATOM	876	CA	ALA	110	-37.313	89.908	-31.160	1.00	26.36
	ATOM	877	CB	ALA	110	-37.973	88.646	-31.664	1.00	20.26
	ATOM	878	C	ALA	110	-36.112	90.237	-32.029	1.00	29.61
	ATOM	879	O	ALA	110	-36.210	91.035	-32.950	1.00	27.61
25	ATOM	880	N	LYS	111	-34.967	89.642	-31.729	1.00	33.77
	ATOM	881	CA	LYS	111	-33.801	89.889	-32.545	1.00	35.09
	ATOM	882	CB	LYS	111	-32.529	89.937	-31.707	1.00	39.38
	ATOM	883	CG	LYS	111	-31.296	90.034	-32.595	1.00	49.74
	ATOM	884	CD	LYS	111	-30.193	90.858	-31.989	1.00	53.95
30	ATOM	885	CE	LYS	111	-29.073	91.053	-33.001	1.00	54.28
	ATOM	886	NZ	LYS	111	-27.949	91.852	-32.405	1.00	60.85
	ATOM	887	C	LYS	111	-33.705	88.776	-33.581	1.00	35.28
	ATOM	888	O	LYS	111	-33.719	87.599	-33.242	1.00	36.58
	ATOM	889	N	LEU	112	-33.620	89.166	-34.846	1.00	32.32
35	ATOM	890	CA	LEU	112	-33.543	88.218	-35.941	1.00	29.67
	ATOM	891	CB	LEU	112	-34.742	88.400	-36.868	1.00	27.04
	ATOM	892	CG	LEU	112	-36.114	88.440	-36.198	1.00	24.38
	ATOM	893	CD1	LEU	112	-37.178	88.763	-37.227	1.00	27.56
	ATOM	894	CD2	LEU	112	-36.396	87.121	-35.533	1.00	19.52
40	ATOM	895	C	LEU	112	-32.257	88.426	-36.726	1.00	32.58
	ATOM	896	O	LEU	112	-31.594	89.453	-36.586	1.00	31.51
	ATOM	897	N	GLU	113	-31.911	87.452	-37.559	1.00	35.75
	ATOM	898	CA	GLU	113	-30.704	87.534	-38.362	1.00	40.08
	ATOM	899	CB	GLU	113	-29.646	86.574	-37.849	1.00	44.23
45	ATOM	900	CG	GLU	113	-29.412	86.591	-36.367	1.00	50.12
	ATOM	901	CD	GLU	113	-28.049	86.027	-36.037	1.00	58.08
	ATOM	902	OE1	GLU	113	-27.631	85.039	-36.710	1.00	61.13
	ATOM	903	OE2	GLU	113	-27.400	86.573	-35.108	1.00	63.64
	ATOM	904	C	GLU	113	-30.960	87.176	-39.804	1.00	41.63
50	ATOM	905	O	GLU	113	-31.946	86.525	-40.124	1.00	48.29
	ATOM	906	N	GLU	114	-30.045	87.601	-40.670	1.00	40.97
	ATOM	907	CA	GLU	114	-30.113	87.305	-42.095	1.00	36.68
	ATOM	908	CB	GLU	114	-28.780	87.595	-42.752	1.00	41.19
	ATOM	909	CG	GLU	114	-28.637	88.918	-43.415	1.00	46.58
55	ATOM	910	CD	GLU	114	-27.604	88.842	-44.519	1.00	51.12
	ATOM	911	OE1	GLU	114	-27.868	88.132	-45.520	1.00	51.88
	ATOM	912	OE2	GLU	114	-26.529	89.475	-44.378	1.00	57.08
	ATOM	913	C	GLU	114	-30.360	85.821	-42.272	1.00	35.51
	ATOM	914	O	GLU	114	-29.581	85.007	-41.790	1.00	38.92
60	ATOM	915	N	GLY	115	-31.424	85.464	-42.972	1.00	32.99
	ATOM	916	CA	GLY	115	-31.681	84.058	-43.189	1.00	31.79
	ATOM	917	C	GLY	115	-32.744	83.483	-42.293	1.00	34.39
	ATOM	918	O	GLY	115	-33.261	82.410	-42.574	1.00	39.20
	ATOM	919	N	ASP	116	-33.056	84.171	-41.203	1.00	35.66
65	ATOM	920	CA	ASP	116	-34.096	83.696	-40.306	1.00	38.09
	ATOM	921	CB	ASP	116	-34.171	84.552	-39.036	1.00	36.00

	ATOM	922	CG	ASP	116	-33.072	84.241	-38.045	1.00	35.09
	ATOM	923	OD1	ASP	116	-32.417	83.185	-38.172	1.00	33.81
	ATOM	924	OD2	ASP	116	-32.874	85.049	-37.120	1.00	35.24
	ATOM	925	C	ASP	116	-35.420	83.803	-41.048	1.00	39.38
5	ATOM	926	O	ASP	116	-35.574	84.633	-41.948	1.00	37.91
	ATOM	927	N	GLU	117	-36.373	82.954	-40.684	1.00	41.54
	ATOM	928	CA	GLU	117	-37.689	83.003	-41.305	1.00	37.92
	ATOM	929	CB	GLU	117	-37.943	81.758	-42.146	1.00	38.90
	ATOM	930	CG	GLU	117	-36.928	81.539	-43.239	1.00	48.15
10	ATOM	931	CD	GLU	117	-37.244	80.315	-44.073	1.00	52.62
	ATOM	932	OE1	GLU	117	-37.715	79.297	-43.496	1.00	55.49
	ATOM	933	OE2	GLU	117	-37.012	80.374	-45.306	1.00	57.10
	ATOM	934	C	GLU	117	-38.726	83.083	-40.201	1.00	34.02
	ATOM	935	O	GLU	117	-38.547	82.497	-39.135	1.00	33.31
15	ATOM	936	N	LEU	118	-39.793	83.831	-40.448	1.00	30.68
	ATOM	937	CA	LEU	118	-40.873	83.961	-39.484	1.00	27.14
	ATOM	938	CB	LEU	118	-41.198	85.434	-39.237	1.00	26.49
	ATOM	939	CG	LEU	118	-40.143	86.307	-38.567	1.00	26.21
	ATOM	940	CD1	LEU	118	-40.641	87.732	-38.502	1.00	23.05
20	ATOM	941	CD2	LEU	118	-39.843	85.784	-37.178	1.00	25.38
	ATOM	942	C	LEU	118	-42.107	83.274	-40.055	1.00	28.33
	ATOM	943	O	LEU	118	-42.328	83.294	-41.267	1.00	31.46
	ATOM	944	N	GLN	119	-42.905	82.652	-39.193	1.00	28.96
	ATOM	945	CA	GLN	119	-44.129	82.007	-39.652	1.00	29.13
25	ATOM	946	CB	GLN	119	-43.844	80.573	-40.074	1.00	28.23
	ATOM	947	CG	GLN	119	-43.514	79.654	-38.937	1.00	36.38
	ATOM	948	CD	GLN	119	-43.173	78.261	-39.409	1.00	36.57
	ATOM	949	OE1	GLN	119	-43.159	77.319	-38.618	1.00	40.70
	ATOM	950	NE2	GLN	119	-42.886	78.121	-40.698	1.00	36.80
30	ATOM	951	C	GLN	119	-45.224	82.038	-38.590	1.00	29.43
	ATOM	952	O	GLN	119	-44.944	82.111	-37.396	1.00	28.80
	ATOM	953	N	LEU	120	-46.470	82.002	-39.051	1.00	32.26
	ATOM	954	CA	LEU	120	-47.652	82.025	-38.194	1.00	29.96
	ATOM	955	CB	LEU	120	-48.643	83.073	-38.720	1.00	31.17
35	ATOM	956	CG	LEU	120	-49.872	83.586	-37.958	1.00	30.26
	ATOM	957	CD1	LEU	120	-50.663	82.447	-37.366	1.00	27.10
	ATOM	958	CD2	LEU	120	-49.415	84.535	-36.887	1.00	30.96
	ATOM	959	C	LEU	120	-48.276	80.630	-38.270	1.00	30.51
	ATOM	960	O	LEU	120	-48.767	80.213	-39.321	1.00	32.88
40	ATOM	961	N	ALA	121	-48.262	79.913	-37.153	1.00	27.40
	ATOM	962	CA	ALA	121	-48.807	78.562	-37.110	1.00	25.55
	ATOM	963	CB	ALA	121	-47.691	77.579	-36.797	1.00	19.41
	ATOM	964	C	ALA	121	-49.952	78.375	-36.115	1.00	29.89
	ATOM	965	O	ALA	121	-49.915	78.892	-35.000	1.00	35.88
45	ATOM	966	N	ILE	122	-50.973	77.634	-36.536	1.00	30.33
	ATOM	967	CA	ILE	122	-52.116	77.338	-35.684	1.00	30.27
	ATOM	968	CB	ILE	122	-53.442	77.574	-36.415	1.00	29.27
	ATOM	969	CG2	ILE	122	-54.590	77.343	-35.460	1.00	30.38
	ATOM	970	CG1	ILE	122	-53.488	79.004	-36.961	1.00	26.03
50	ATOM	971	CD1	ILE	122	-54.747	79.346	-37.737	1.00	29.21
	ATOM	972	C	ILE	122	-51.996	75.861	-35.323	1.00	33.20
	ATOM	973	O	ILE	122	-52.089	74.997	-36.192	1.00	33.30
	ATOM	974	N	PRO	123	-51.779	75.555	-34.032	1.00	34.70
	ATOM	975	CD	PRO	123	-51.616	76.525	-32.945	1.00	34.40
55	ATOM	976	CA	PRO	123	-51.630	74.188	-33.516	1.00	36.49
	ATOM	977	CB	PRO	123	-51.076	74.400	-32.104	1.00	30.22
	ATOM	978	CG	PRO	123	-50.611	75.828	-32.090	1.00	36.72
	ATOM	979	C	PRO	123	-52.929	73.385	-33.476	1.00	39.64
	ATOM	980	O	PRO	123	-53.324	72.898	-32.413	1.00	40.48
60	ATOM	981	N	ARG	124	-53.586	73.249	-34.625	1.00	44.58
	ATOM	982	CA	ARG	124	-54.835	72.505	-34.707	1.00	48.04
	ATOM	983	CB	ARG	124	-56.018	73.426	-34.435	1.00	52.91
	ATOM	984	CG	ARG	124	-55.923	74.175	-33.118	1.00	60.50
	ATOM	985	CD	ARG	124	-56.943	73.607	-32.156	1.00	72.57
65	ATOM	986	NE	ARG	124	-56.801	74.172	-30.803	1.00	84.90
	ATOM	987	CZ	ARG	124	-56.049	73.654	-29.823	1.00	85.44

	ATOM	988	NH1	ARG	124	-55.349	72.538	-30.033	1.00	87.69
	ATOM	989	NH2	ARG	124	-55.995	74.259	-28.629	1.00	82.78
	ATOM	990	C	ARG	124	-55.004	71.836	-36.061	1.00	48.39
	ATOM	991	O	ARG	124	-54.551	72.358	-37.080	1.00	45.26
5	ATOM	992	N	GLU	125	-55.670	70.683	-36.048	1.00	52.84
	ATOM	993	CA	GLU	125	-55.913	69.878	-37.242	1.00	55.02
	ATOM	994	CB	GLU	125	-56.852	68.719	-36.896	1.00	63.62
	ATOM	995	CG	GLU	125	-57.827	69.025	-35.756	1.00	70.62
	ATOM	996	CD	GLU	125	-57.118	69.487	-34.488	1.00	74.76
10	ATOM	997	OE1	GLU	125	-56.364	68.673	-33.891	1.00	75.75
	ATOM	998	OE2	GLU	125	-57.316	70.663	-34.093	1.00	80.71
	ATOM	999	C	GLU	125	-56.455	70.642	-38.442	1.00	53.44
	ATOM	1000	O	GLU	125	-55.747	70.815	-39.435	1.00	57.50
	ATOM	1001	N	ASN	126	-57.711	71.061	-38.388	1.00	47.18
15	ATOM	1002	CA	ASN	126	-58.264	71.828	-39.497	1.00	47.59
	ATOM	1003	CB	ASN	126	-59.265	71.011	-40.310	1.00	51.03
	ATOM	1004	CG	ASN	126	-58.663	70.480	-41.602	1.00	57.07
	ATOM	1005	OD1	ASN	126	-57.882	69.515	-41.601	1.00	61.15
	ATOM	1006	ND2	ASN	126	-59.008	71.124	-42.718	1.00	56.63
20	ATOM	1007	C	ASN	126	-58.930	73.048	-38.922	1.00	47.06
	ATOM	1008	O	ASN	126	-60.158	73.177	-38.920	1.00	45.61
	ATOM	1009	N	ALA	127	-58.088	73.941	-38.420	1.00	43.50
	ATOM	1010	CA	ALA	127	-58.538	75.164	-37.795	1.00	38.26
	ATOM	1011	CB	ALA	127	-57.359	76.085	-37.584	1.00	37.97
25	ATOM	1012	C	ALA	127	-59.609	75.869	-38.604	1.00	37.16
	ATOM	1013	O	ALA	127	-59.447	76.106	-39.800	1.00	33.14
	ATOM	1014	N	GLN	128	-60.718	76.179	-37.944	1.00	37.87
	ATOM	1015	CA	GLN	128	-61.797	76.912	-38.589	1.00	37.19
	ATOM	1016	CB	GLN	128	-63.127	76.563	-37.929	1.00	40.23
30	ATOM	1017	CG	GLN	128	-63.541	75.137	-38.206	1.00	38.14
	ATOM	1018	CD	GLN	128	-63.563	74.854	-39.690	1.00	39.51
	ATOM	1019	OE1	GLN	128	-64.397	75.389	-40.427	1.00	40.36
	ATOM	1020	NE2	GLN	128	-62.626	74.024	-40.148	1.00	41.53
	ATOM	1021	C	GLN	128	-61.433	78.381	-38.394	1.00	35.55
35	ATOM	1022	O	GLN	128	-61.640	78.969	-37.329	1.00	35.57
	ATOM	1023	N	ILE	129	-60.873	78.953	-39.449	1.00	34.64
	ATOM	1024	CA	ILE	129	-60.367	80.318	-39.450	1.00	29.60
	ATOM	1025	CB	ILE	129	-58.808	80.242	-39.642	1.00	31.03
	ATOM	1026	CG2	ILE	129	-58.311	81.207	-40.687	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-58.129	80.436	-38.305	1.00	26.37
	ATOM	1028	CD1	ILE	129	-58.499	79.394	-37.318	1.00	34.70
	ATOM	1029	C	ILE	129	-61.002	81.191	-40.528	1.00	29.55
	ATOM	1030	O	ILE	129	-61.471	80.690	-41.539	1.00	33.42
	ATOM	1031	N	SER	130	-61.027	82.500	-40.307	1.00	27.67
45	ATOM	1032	CA	SER	130	-61.561	83.417	-41.309	1.00	25.00
	ATOM	1033	CB	SER	130	-62.207	84.628	-40.658	1.00	22.09
	ATOM	1034	OG	SER	130	-62.503	85.607	-41.644	1.00	20.89
	ATOM	1035	C	SER	130	-60.399	83.890	-42.168	1.00	26.03
	ATOM	1036	O	SER	130	-59.349	84.216	-41.645	1.00	30.77
50	ATOM	1037	N	LEU	131	-60.581	83.936	-43.479	1.00	24.49
	ATOM	1038	CA	LEU	131	-59.507	84.363	-44.362	1.00	28.07
	ATOM	1039	CB	LEU	131	-59.307	83.343	-45.483	1.00	28.87
	ATOM	1040	CG	LEU	131	-58.347	82.190	-45.192	1.00	30.76
	ATOM	1041	CD1	LEU	131	-58.634	81.570	-43.857	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-58.475	81.165	-46.289	1.00	33.37
	ATOM	1043	C	LEU	131	-59.687	85.747	-44.961	1.00	33.66
	ATOM	1044	O	LEU	131	-59.227	86.010	-46.073	1.00	37.67
	ATOM	1045	N	ASP	132	-60.355	86.631	-44.228	1.00	34.97
	ATOM	1046	CA	ASP	132	-60.556	88.000	-44.692	1.00	34.31
60	ATOM	1047	CB	ASP	132	-61.780	88.627	-44.025	1.00	42.68
	ATOM	1048	CG	ASP	132	-63.089	88.127	-44.614	1.00	47.98
	ATOM	1049	OD1	ASP	132	-64.156	88.407	-44.014	1.00	50.41
	ATOM	1050	OD2	ASP	132	-63.045	87.465	-45.681	1.00	46.42
	ATOM	1051	C	ASP	132	-59.312	88.805	-44.358	1.00	34.32
65	ATOM	1052	O	ASP	132	-58.796	88.737	-43.248	1.00	32.87
	ATOM	1053	N	GLY	133	-58.836	89.566	-45.335	1.00	34.53

	ATOM	1054	CA	GLY	133	-57.636	90.356	-45.149	1.00	36.93
	ATOM	1055	C	GLY	133	-57.603	91.304	-43.966	1.00	35.06
	ATOM	1056	O	GLY	133	-56.526	91.686	-43.509	1.00	40.44
	ATOM	1057	N	ASP	134	-58.765	91.686	-43.457	1.00	31.08
5	ATOM	1058	CA	ASP	134	-58.793	92.612	-42.342	1.00	27.63
	ATOM	1059	CB	ASP	134	-59.961	93.589	-42.501	1.00	28.43
	ATOM	1060	CG	ASP	134	-61.308	92.902	-42.542	1.00	31.78
	ATOM	1061	OD1	ASP	134	-61.392	91.777	-43.069	1.00	35.68
	ATOM	1062	OD2	ASP	134	-62.294	93.498	-42.061	1.00	32.53
10	ATOM	1063	C	ASP	134	-58.815	91.947	-40.978	1.00	25.08
	ATOM	1064	O	ASP	134	-58.442	92.554	-40.000	1.00	19.71
	ATOM	1065	N	VAL	135	-59.211	90.686	-40.905	1.00	26.84
	ATOM	1066	CA	VAL	135	-59.245	90.026	-39.608	1.00	23.72
	ATOM	1067	CB	VAL	135	-60.536	89.227	-39.423	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-61.688	90.182	-39.271	1.00	22.76
	ATOM	1069	CG2	VAL	135	-60.757	88.302	-40.601	1.00	22.12
	ATOM	1070	C	VAL	135	-58.045	89.139	-39.292	1.00	26.54
	ATOM	1071	O	VAL	135	-57.744	88.919	-38.126	1.00	29.26
	ATOM	1072	N	THR	136	-57.361	88.617	-40.308	1.00	26.39
20	ATOM	1073	CA	THR	136	-56.175	87.807	-40.034	1.00	28.40
	ATOM	1074	CB	THR	136	-56.461	86.275	-40.185	1.00	28.21
	ATOM	1075	OG1	THR	136	-55.929	85.792	-41.417	1.00	37.13
	ATOM	1076	CG2	THR	136	-57.943	85.999	-40.137	1.00	26.90
	ATOM	1077	C	THR	136	-54.998	88.257	-40.911	1.00	31.60
25	ATOM	1078	O	THR	136	-55.047	88.187	-42.142	1.00	27.29
	ATOM	1079	N	PHE	137	-53.947	88.744	-40.249	1.00	31.45
	ATOM	1080	CA	PHE	137	-52.765	89.259	-40.929	1.00	26.64
	ATOM	1081	CB	PHE	137	-52.935	90.757	-41.134	1.00	26.84
	ATOM	1082	CG	PHE	137	-53.497	91.471	-39.939	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-52.661	91.920	-38.921	1.00	27.77
	ATOM	1084	CD2	PHE	137	-54.858	91.704	-39.832	1.00	23.70
	ATOM	1085	CE1	PHE	137	-53.172	92.590	-37.821	1.00	27.32
	ATOM	1086	CE2	PHE	137	-55.376	92.373	-38.734	1.00	26.31
	ATOM	1087	CZ	PHE	137	-54.529	92.817	-37.729	1.00	29.43
35	ATOM	1088	C	PHE	137	-51.473	88.947	-40.177	1.00	28.06
	ATOM	1089	O	PHE	137	-51.512	88.572	-39.014	1.00	25.04
	ATOM	1090	N	PHE	138	-50.333	89.111	-40.847	1.00	29.81
	ATOM	1091	CA	PHE	138	-49.036	88.782	-40.260	1.00	29.93
	ATOM	1092	CB	PHE	138	-48.542	87.496	-40.920	1.00	27.47
40	ATOM	1093	CG	PHE	138	-47.376	86.863	-40.241	1.00	28.49
	ATOM	1094	CD1	PHE	138	-47.111	87.103	-38.897	1.00	27.65
	ATOM	1095	CD2	PHE	138	-46.535	86.011	-40.952	1.00	31.06
	ATOM	1096	CE1	PHE	138	-46.024	86.505	-38.274	1.00	30.90
	ATOM	1097	CE2	PHE	138	-45.444	85.406	-40.342	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-45.186	85.653	-39.001	1.00	27.41
	ATOM	1099	C	PHE	138	-47.998	89.918	-40.353	1.00	30.91
	ATOM	1100	O	PHE	138	-47.734	90.459	-41.421	1.00	24.94
	ATOM	1101	N	GLY	139	-47.407	90.220	-39.194	1.00	36.94
	ATOM	1102	CA	GLY	139	-46.459	91.311	-38.973	1.00	36.08
50	ATOM	1103	C	GLY	139	-45.102	91.566	-39.580	1.00	36.48
	ATOM	1104	O	GLY	139	-44.953	91.524	-40.787	1.00	43.61
	ATOM	1105	N	ALA	140	-44.139	91.907	-38.722	1.00	36.60
	ATOM	1106	CA	ALA	140	-42.742	92.212	-39.088	1.00	34.94
	ATOM	1107	CB	ALA	140	-42.240	91.218	-40.120	1.00	36.94
55	ATOM	1108	C	ALA	140	-42.404	93.643	-39.550	1.00	32.53
	ATOM	1109	O	ALA	140	-42.693	94.040	-40.673	1.00	27.96
	ATOM	1110	N	LEU	141	-41.759	94.395	-38.662	1.00	33.23
	ATOM	1111	CA	LEU	141	-41.332	95.778	-38.917	1.00	36.11
	ATOM	1112	CB	LEU	141	-42.319	96.750	-38.270	1.00	37.53
60	ATOM	1113	CG	LEU	141	-42.075	98.262	-38.299	1.00	38.87
	ATOM	1114	CD1	LEU	141	-43.325	98.963	-37.825	1.00	40.67
	ATOM	1115	CD2	LEU	141	-40.911	98.653	-37.410	1.00	37.53
	ATOM	1116	C	LEU	141	-39.940	95.972	-38.307	1.00	38.89
	ATOM	1117	O	LEU	141	-39.724	95.656	-37.139	1.00	39.25
65	ATOM	1118	N	LYS	142	-38.993	96.497	-39.076	1.00	42.86
	ATOM	1119	CA	LYS	142	-37.650	96.679	-38.535	1.00	42.96

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	ATOM	1120	CB	LYS	142	-36.614	96.614	-39.653	1.00	41.80
	ATOM	1121	CG	LYS	142	-35.209	96.827	-39.142	1.00	46.88
	ATOM	1122	CD	LYS	142	-34.154	96.310	-40.098	1.00	49.58
	ATOM	1123	CE	LYS	142	-32.773	96.557	-39.512	1.00	51.07
5	ATOM	1124	NZ	LYS	142	-31.690	96.000	-40.359	1.00	56.20
	ATOM	1125	C	LYS	142	-37.447	97.956	-37.727	1.00	43.41
	ATOM	1126	O	LYS	142	-37.777	99.049	-38.179	1.00	44.98
	ATOM	1127	N	LEU	143	-36.896	97.808	-36.526	1.00	42.81
	ATOM	1128	CA	LEU	143	-36.639	98.948	-35.648	1.00	40.65
10	ATOM	1129	CB	LEU	143	-36.603	98.497	-34.189	1.00	36.56
	ATOM	1130	CG	LEU	143	-37.828	97.782	-33.619	1.00	36.17
	ATOM	1131	CD1	LEU	143	-37.543	97.375	-32.185	1.00	37.95
	ATOM	1132	CD2	LEU	143	-39.037	98.689	-33.685	1.00	27.38
	ATOM	1133	C	LEU	143	-35.299	99.591	-35.988	1.00	43.14
15	ATOM	1134	O	LEU	143	-34.367	98.909	-36.426	1.00	46.28
	ATOM	1135	N	LEU	144	-35.196	100.898	-35.786	1.00	42.63
	ATOM	1136	CA	LEU	144	-33.948	101.592	-36.065	1.00	44.11
	ATOM	1137	CB	LEU	144	-34.192	103.089	-36.235	1.00	44.88
	ATOM	1138	CG	LEU	144	-35.047	103.456	-37.446	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-35.308	104.938	-37.460	1.00	47.71
	ATOM	1140	CD2	LEU	144	-34.339	103.032	-38.720	1.00	47.87
	ATOM	1141	C	LEU	144	-32.971	101.372	-34.927	1.00	45.66
	ATOM	1142	O	LEU	144	-33.423	101.026	-33.820	1.00	46.37
	ATOM	1143	OXT	LEU	144	-31.762	101.562	-35.157	1.00	50.66
25	END					0.000	0.000	0.000	0.00	0.00

TABLE 10

	11								
	ATOM	1	CB	VAL	1	-28.702	106.858	-34.286	1.00 69.28
5	ATOM	2	CG1	VAL	1	-28.176	106.176	-35.569	1.00 68.26
	ATOM	3	CG2	VAL	1	-27.981	108.187	-34.030	1.00 71.75
	ATOM	4	C	VAL	1	-30.944	105.728	-34.247	1.00 64.86
	ATOM	5	O	VAL	1	-31.422	105.154	-35.237	1.00 64.02
	ATOM	6	N	VAL	1	-30.722	108.050	-33.350	1.00 65.10
10	ATOM	7	CA	VAL	1	-30.243	107.092	-34.393	1.00 65.98
	ATOM	8	N	THR	2	-30.992	105.208	-33.017	1.00 61.42
	ATOM	9	CA	THR	2	-31.636	103.920	-32.761	1.00 56.64
	ATOM	10	CB	THR	2	-30.655	102.893	-32.170	1.00 56.77
	ATOM	11	OG1	THR	2	-30.059	103.441	-30.991	1.00 57.45
15	ATOM	12	CG2	THR	2	-29.566	102.535	-33.174	1.00 55.68
	ATOM	13	C	THR	2	-32.783	104.054	-31.778	1.00 53.08
	ATOM	14	O	THR	2	-32.957	105.089	-31.143	1.00 52.00
	ATOM	15	N	GLN	3	-33.556	102.984	-31.643	1.00 50.34
	ATOM	16	CA	GLN	3	-34.692	102.973	-30.731	1.00 45.97
20	ATOM	17	CB	GLN	3	-35.951	102.533	-31.470	1.00 46.09
	ATOM	18	CG	GLN	3	-36.178	103.256	-32.772	1.00 49.60
	ATOM	19	CD	GLN	3	-37.473	102.847	-33.426	1.00 50.43
	ATOM	20	OE1	GLN	3	-38.548	103.140	-32.912	1.00 50.94
	ATOM	21	NE2	GLN	3	-37.380	102.157	-34.561	1.00 49.99
25	ATOM	22	C	GLN	3	-34.444	102.032	-29.567	1.00 42.41
	ATOM	23	O	GLN	3	-34.426	100.815	-29.726	1.00 37.34
	ATOM	24	N	ASP	4	-34.250	102.603	-28.390	1.00 40.32
	ATOM	25	CA	ASP	4	-34.006	101.802	-27.204	1.00 39.37
	ATOM	26	CB	ASP	4	-33.599	102.701	-26.034	1.00 44.56
30	ATOM	27	CG	ASP	4	-32.268	103.387	-26.264	1.00 45.76
	ATOM	28	OD1	ASP	4	-31.842	103.465	-27.439	1.00 48.58
	ATOM	29	OD2	ASP	4	-31.658	103.852	-25.278	1.00 45.55
	ATOM	30	C	ASP	4	-35.269	101.048	-26.848	1.00 36.30
	ATOM	31	O	ASP	4	-36.376	101.534	-27.066	1.00 36.29
35	ATOM	32	N	CYS	5	-35.098	99.851	-26.311	1.00 35.11
	ATOM	33	CA	CYS	5	-36.229	99.043	-25.892	1.00 35.61
	ATOM	34	CB	CYS	5	-36.924	98.399	-27.094	1.00 34.96
	ATOM	35	SG	CYS	5	-35.834	97.728	-28.334	1.00 35.34
	ATOM	36	C	CYS	5	-35.766	97.990	-24.908	1.00 31.80
40	ATOM	37	O	CYS	5	-34.617	97.566	-24.933	1.00 31.85
	ATOM	38	N	LEU	6	-36.661	97.603	-24.014	1.00 27.72
	ATOM	39	CA	LEU	6	-36.367	96.596	-23.016	1.00 30.80
	ATOM	40	CB	LEU	6	-36.069	97.254	-21.674	1.00 29.25
	ATOM	41	CG	LEU	6	-35.809	96.340	-20.476	1.00 29.47
45	ATOM	42	CD1	LEU	6	-34.966	97.064	-19.459	1.00 30.65
	ATOM	43	CD2	LEU	6	-37.111	95.929	-19.857	1.00 35.27
	ATOM	44	C	LEU	6	-37.587	95.703	-22.909	1.00 31.83
	ATOM	45	O	LEU	6	-38.697	96.192	-22.734	1.00 33.45
	ATOM	46	N	GLN	7	-37.388	94.395	-23.027	1.00 32.23
50	ATOM	47	CA	GLN	7	-38.494	93.465	-22.954	1.00 30.26
	ATOM	48	CB	GLN	7	-38.605	92.698	-24.266	1.00 29.56
	ATOM	49	CG	GLN	7	-39.885	91.911	-24.439	1.00 25.09
	ATOM	50	CD	GLN	7	-40.038	91.392	-25.855	1.00 26.50
	ATOM	51	OE1	GLN	7	-39.323	90.486	-26.284	1.00 24.40
55	ATOM	52	NE2	GLN	7	-40.960	91.980	-26.595	1.00 27.03
	ATOM	53	C	GLN	7	-38.301	92.509	-21.797	1.00 32.11
	ATOM	54	O	GLN	7	-37.193	92.065	-21.522	1.00 34.24
	ATOM	55	N	LEU	8	-39.397	92.215	-21.114	1.00 33.73
	ATOM	56	CA	LEU	8	-39.391	91.312	-19.974	1.00 35.41
60	ATOM	57	CB	LEU	8	-39.942	92.031	-18.731	1.00 32.44
	ATOM	58	CG	LEU	8	-38.996	92.737	-17.749	1.00 27.69
	ATOM	59	CD1	LEU	8	-37.651	93.016	-18.370	1.00 26.43
	ATOM	60	CD2	LEU	8	-39.659	94.002	-17.277	1.00 18.64
	ATOM	61	C	LEU	8	-40.223	90.076	-20.281	1.00 36.87
65	ATOM	62	O	LEU	8	-41.216	90.136	-21.005	1.00 36.05
	ATOM	63	N	ILE	9	-39.807	88.955	-19.711	1.00 39.25

	ATOM	64	CA	ILE	9	-40.469	87.677	-19.912	1.00	35.57
	ATOM	65	CB	ILE	9	-39.522	86.751	-20.717	1.00	34.28
	ATOM	66	CG2	ILE	9	-39.454	85.367	-20.120	1.00	38.20
	ATOM	67	CG1	ILE	9	-39.975	86.694	-22.157	1.00	35.22
5	ATOM	68	CD1	ILE	9	-39.129	85.766	-22.961	1.00	43.63
	ATOM	69	C	ILE	9	-40.826	87.066	-18.554	1.00	35.28
	ATOM	70	O	ILE	9	-40.111	87.260	-17.576	1.00	33.75
	ATOM	71	N	ALA	10	-41.937	86.343	-18.486	1.00	35.78
	ATOM	72	CA	ALA	10	-42.331	85.723	-17.230	1.00	35.28
10	ATOM	73	CB	ALA	10	-43.678	85.054	-17.375	1.00	28.47
	ATOM	74	C	ALA	10	-41.290	84.702	-16.790	1.00	36.49
	ATOM	75	O	ALA	10	-40.757	83.950	-17.601	1.00	37.41
	ATOM	76	N	ASP	11	-41.002	84.682	-15.497	1.00	39.44
	ATOM	77	CA	ASP	11	-40.030	83.748	-14.942	1.00	40.77
15	ATOM	78	CB	ASP	11	-39.079	84.482	-13.996	1.00	38.90
	ATOM	79	CG	ASP	11	-38.131	83.545	-13.289	1.00	40.34
	ATOM	80	OD1	ASP	11	-37.848	82.450	-13.834	1.00	39.50
	ATOM	81	OD2	ASP	11	-37.664	83.910	-12.194	1.00	37.38
	ATOM	82	C	ASP	11	-40.748	82.625	-14.201	1.00	42.25
20	ATOM	83	O	ASP	11	-41.077	82.746	-13.020	1.00	40.54
	ATOM	84	N	SER	12	-40.980	81.529	-14.913	1.00	42.92
	ATOM	85	CA	SER	12	-41.692	80.373	-14.371	1.00	43.84
	ATOM	86	CB	SER	12	-41.916	79.348	-15.476	1.00	42.13
	ATOM	87	OG	SER	12	-40.679	78.948	-16.038	1.00	43.31
25	ATOM	88	C	SER	12	-41.015	79.688	-13.196	1.00	46.10
	ATOM	89	O	SER	12	-41.573	78.761	-12.608	1.00	43.90
	ATOM	90	N	GLU	13	-39.818	80.145	-12.848	1.00	46.72
	ATOM	91	CA	GLU	13	-39.091	79.540	-11.748	1.00	44.93
	ATOM	92	CB	GLU	13	-37.647	79.307	-12.153	1.00	46.71
30	ATOM	93	CG	GLU	13	-37.533	78.267	-13.233	1.00	58.25
	ATOM	94	CD	GLU	13	-36.175	77.594	-13.239	1.00	66.70
	ATOM	95	OE1	GLU	13	-35.808	77.011	-12.181	1.00	74.10
	ATOM	96	OE2	GLU	13	-35.478	77.643	-14.288	1.00	71.55
	ATOM	97	C	GLU	13	-39.160	80.272	-10.412	1.00	43.24
35	ATOM	98	O	GLU	13	-38.390	79.976	-9.496	1.00	44.25
	ATOM	99	N	THR	14	-40.066	81.232	-10.298	1.00	37.06
	ATOM	100	CA	THR	14	-40.244	81.925	-9.027	1.00	36.77
	ATOM	101	CB	THR	14	-39.471	83.276	-8.955	1.00	33.39
	ATOM	102	OG1	THR	14	-40.134	84.262	-9.744	1.00	41.69
40	ATOM	103	CG2	THR	14	-38.046	83.105	-9.456	1.00	31.66
	ATOM	104	C	THR	14	-41.743	82.160	-8.873	1.00	34.96
	ATOM	105	O	THR	14	-42.462	82.293	-9.856	1.00	35.04
	ATOM	106	N	PRO	15	-42.232	82.189	-7.634	1.00	37.11
	ATOM	107	CD	PRO	15	-41.464	82.010	-6.393	1.00	36.50
45	ATOM	108	CA	PRO	15	-43.655	82.397	-7.342	1.00	38.81
	ATOM	109	CB	PRO	15	-43.726	82.225	-5.825	1.00	39.15
	ATOM	110	CG	PRO	15	-42.496	81.416	-5.492	1.00	37.32
	ATOM	111	C	PRO	15	-44.154	83.772	-7.765	1.00	37.78
	ATOM	112	O	PRO	15	-43.383	84.727	-7.795	1.00	35.91
50	ATOM	113	N	THR	16	-45.439	83.882	-8.087	1.00	36.67
	ATOM	114	CA	THR	16	-45.976	85.178	-8.470	1.00	38.46
	ATOM	115	CB	THR	16	-47.426	85.078	-8.979	1.00	38.61
	ATOM	116	OG1	THR	16	-48.276	84.616	-7.924	1.00	42.44
	ATOM	117	CG2	THR	16	-47.510	84.115	-10.152	1.00	40.11
55	ATOM	118	C	THR	16	-45.963	86.016	-7.209	1.00	35.97
	ATOM	119	O	THR	16	-46.298	85.520	-6.142	1.00	39.85
	ATOM	120	N	ILE	17	-45.572	87.278	-7.327	1.00	33.66
	ATOM	121	CA	ILE	17	-45.514	88.159	-6.170	1.00	34.76
	ATOM	122	CB	ILE	17	-44.639	89.384	-6.465	1.00	32.44
60	ATOM	123	CG2	ILE	17	-44.685	90.351	-5.298	1.00	28.66
	ATOM	124	CG1	ILE	17	-43.204	88.927	-6.756	1.00	33.15
	ATOM	125	CD1	ILE	17	-42.256	90.045	-7.141	1.00	31.19
	ATOM	126	C	ILE	17	-46.882	88.637	-5.703	1.00	40.64
	ATOM	127	O	ILE	17	-47.667	89.158	-6.485	1.00	44.07
65	ATOM	128	N	GLN	18	-47.167	88.456	-4.416	1.00	45.91
	ATOM	129	CA	GLN	18	-48.441	88.885	-3.844	1.00	50.22

	ATOM	130	CB	GLN	18	-49.030	87.795	-2.955	1.00	53.09
	ATOM	131	CG	GLN	18	-50.411	87.377	-3.385	1.00	57.60
	ATOM	132	CD	GLN	18	-50.371	86.655	-4.706	1.00	61.39
	ATOM	133	OE1	GLN	18	-49.866	85.533	-4.798	1.00	68.36
5	ATOM	134	NE2	GLN	18	-50.884	87.297	-5.747	1.00	64.93
	ATOM	135	C	GLN	18	-48.258	90.142	-3.012	1.00	51.56
	ATOM	136	O	GLN	18	-47.355	90.223	-2.185	1.00	56.44
	ATOM	137	N	LYS	19	-49.121	91.121	-3.222	1.00	52.60
	ATOM	138	CA	LYS	19	-49.018	92.357	-2.465	1.00	56.10
10	ATOM	139	CB	LYS	19	-47.771	93.139	-2.881	1.00	55.55
	ATOM	140	CG	LYS	19	-47.693	94.481	-2.191	1.00	61.15
	ATOM	141	CD	LYS	19	-46.370	95.186	-2.406	1.00	64.42
	ATOM	142	CE	LYS	19	-46.360	96.529	-1.662	1.00	64.84
	ATOM	143	NZ	LYS	19	-45.061	97.263	-1.805	1.00	69.52
15	ATOM	144	C	LYS	19	-50.244	93.244	-2.621	1.00	57.41
	ATOM	145	O	LYS	19	-50.735	93.444	-3.733	1.00	61.02
	ATOM	146	N	GLY	20	-50.724	93.777	-1.498	1.00	58.84
	ATOM	147	CA	GLY	20	-51.888	94.644	-1.509	1.00	56.66
	ATOM	148	C	GLY	20	-53.049	94.069	-2.297	1.00	56.87
20	ATOM	149	O	GLY	20	-53.756	94.812	-2.979	1.00	58.64
	ATOM	150	N	SER	21	-53.243	92.752	-2.201	1.00	54.49
	ATOM	151	CA	SER	21	-54.321	92.056	-2.907	1.00	53.68
	ATOM	152	CB	SER	21	-55.689	92.585	-2.452	1.00	55.80
	ATOM	153	OG	SER	21	-56.069	93.745	-3.168	1.00	65.34
25	ATOM	154	C	SER	21	-54.190	92.160	-4.446	1.00	51.41
	ATOM	155	O	SER	21	-55.182	92.089	-5.191	1.00	47.74
	ATOM	156	N	TYR	22	-52.950	92.333	-4.903	1.00	46.25
	ATOM	157	CA	TYR	22	-52.628	92.418	-6.320	1.00	40.57
	ATOM	158	CB	TYR	22	-52.000	93.774	-6.651	1.00	43.27
30	ATOM	159	CG	TYR	22	-52.999	94.853	-6.981	1.00	49.18
	ATOM	160	CD1	TYR	22	-53.971	95.236	-6.063	1.00	52.61
	ATOM	161	CE1	TYR	22	-54.907	96.227	-6.369	1.00	51.92
	ATOM	162	CD2	TYR	22	-52.982	95.489	-8.218	1.00	50.01
	ATOM	163	CE2	TYR	22	-53.910	96.482	-8.533	1.00	50.94
35	ATOM	164	CZ	TYR	22	-54.869	96.845	-7.605	1.00	50.78
	ATOM	165	OH	TYR	22	-55.788	97.822	-7.917	1.00	53.87
	ATOM	166	C	TYR	22	-51.624	91.313	-6.632	1.00	37.85
	ATOM	167	O	TYR	22	-50.830	90.933	-5.773	1.00	38.70
	ATOM	168	N	THR	23	-51.666	90.782	-7.847	1.00	33.81
40	ATOM	169	CA	THR	23	-50.717	89.744	-8.234	1.00	32.40
	ATOM	170	CB	THR	23	-51.400	88.536	-8.924	1.00	31.58
	ATOM	171	OG1	THR	23	-52.405	87.983	-8.068	1.00	30.76
	ATOM	172	CG2	THR	23	-50.375	87.468	-9.236	1.00	24.08
	ATOM	173	C	THR	23	-49.742	90.356	-9.224	1.00	33.28
45	ATOM	174	O	THR	23	-50.151	91.018	-10.171	1.00	30.44
	ATOM	175	N	PHE	24	-48.455	90.136	-9.000	1.00	32.16
	ATOM	176	CA	PHE	24	-47.435	90.663	-9.888	1.00	30.68
	ATOM	177	CB	PHE	24	-46.505	91.613	-9.134	1.00	28.20
	ATOM	178	CG	PHE	24	-47.177	92.863	-8.661	1.00	31.00
50	ATOM	179	CD1	PHE	24	-47.880	92.880	-7.468	1.00	31.69
	ATOM	180	CD2	PHE	24	-47.135	94.023	-9.429	1.00	31.42
	ATOM	181	CE1	PHE	24	-48.532	94.032	-7.047	1.00	32.57
	ATOM	182	CE2	PHE	24	-47.782	95.175	-9.016	1.00	25.01
	ATOM	183	CZ	PHE	24	-48.481	95.179	-7.822	1.00	30.12
55	ATOM	184	C	PHE	24	-46.618	89.551	-10.533	1.00	31.73
	ATOM	185	O	PHE	24	-46.126	88.651	-9.859	1.00	33.31
	ATOM	186	N	VAL	25	-46.486	89.616	-11.848	1.00	32.92
	ATOM	187	CA	VAL	25	-45.723	88.628	-12.579	1.00	30.84
	ATOM	188	CB	VAL	25	-45.909	88.802	-14.095	1.00	30.29
60	ATOM	189	CG1	VAL	25	-45.074	87.790	-14.840	1.00	28.13
	ATOM	190	CG2	VAL	25	-47.362	88.659	-14.456	1.00	29.53
	ATOM	191	C	VAL	25	-44.239	88.777	-12.266	1.00	36.08
	ATOM	192	O	VAL	25	-43.717	89.888	-12.208	1.00	35.82
	ATOM	193	N	PRO	26	-43.543	87.655	-12.033	1.00	38.03
65	ATOM	194	CD	PRO	26	-44.116	86.325	-11.775	1.00	38.28
	ATOM	195	CA	PRO	26	-42.105	87.662	-11.731	1.00	38.08

	ATOM	196	CB	PRO	26	-41.865	86.280	-11.118	1.00	37.62
	ATOM	197	CG	PRO	26	-43.222	85.823	-10.688	1.00	36.31
	ATOM	198	C	PRO	26	-41.393	87.796	-13.076	1.00	40.85
	ATOM	199	O	PRO	26	-41.432	86.876	-13.891	1.00	42.27
5	ATOM	200	N	TRP	27	-40.744	88.925	-13.323	1.00	40.86
	ATOM	201	CA	TRP	27	-40.093	89.106	-14.615	1.00	37.05
	ATOM	202	CB	TRP	27	-40.141	90.577	-15.035	1.00	33.29
	ATOM	203	CG	TRP	27	-41.522	91.108	-15.175	1.00	31.06
	ATOM	204	CD2	TRP	27	-42.530	90.647	-16.076	1.00	27.79
10	ATOM	205	CE2	TRP	27	-43.681	91.415	-15.836	1.00	29.28
	ATOM	206	CE3	TRP	27	-42.573	89.660	-17.060	1.00	27.06
	ATOM	207	CD1	TRP	27	-42.082	92.108	-14.452	1.00	31.07
	ATOM	208	NE1	TRP	27	-43.378	92.302	-14.839	1.00	32.70
	ATOM	209	CZ2	TRP	27	-44.871	91.228	-16.545	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-43.757	89.473	-17.766	1.00	26.93
	ATOM	211	CH2	TRP	27	-44.888	90.255	-17.503	1.00	28.08
	ATOM	212	C	TRP	27	-38.666	88.616	-14.706	1.00	38.50
	ATOM	213	O	TRP	27	-37.971	88.473	-13.706	1.00	39.07
	ATOM	214	N	LEU	28	-38.253	88.358	-15.942	1.00	40.19
20	ATOM	215	CA	LEU	28	-36.908	87.907	-16.264	1.00	39.45
	ATOM	216	CB	LEU	28	-36.915	86.421	-16.598	1.00	44.18
	ATOM	217	CG	LEU	28	-35.553	85.730	-16.577	1.00	46.83
	ATOM	218	CD1	LEU	28	-35.037	85.700	-15.136	1.00	45.15
	ATOM	219	CD2	LEU	28	-35.682	84.311	-17.139	1.00	49.15
25	ATOM	220	C	LEU	28	-36.545	88.704	-17.510	1.00	39.80
	ATOM	221	O	LEU	28	-37.379	88.881	-18.386	1.00	42.40
	ATOM	222	N	LEU	29	-35.320	89.194	-17.597	1.00	33.58
	ATOM	223	CA	LEU	29	-34.943	89.965	-18.766	1.00	29.21
	ATOM	224	CB	LEU	29	-33.549	90.567	-18.596	1.00	27.16
30	ATOM	225	CG	LEU	29	-33.050	91.343	-19.819	1.00	22.36
	ATOM	226	CD1	LEU	29	-33.784	92.670	-19.916	1.00	21.47
	ATOM	227	CD2	LEU	29	-31.564	91.565	-19.727	1.00	14.53
	ATOM	228	C	LEU	29	-34.963	89.136	-20.044	1.00	29.98
	ATOM	229	O	LEU	29	-34.352	88.075	-20.127	1.00	30.36
35	ATOM	230	N	SER	30	-35.683	89.636	-21.040	1.00	29.88
	ATOM	231	CA	SER	30	-35.756	88.979	-22.331	1.00	28.49
	ATOM	232	CB	SER	30	-37.085	89.284	-23.022	1.00	26.13
	ATOM	233	OG	SER	30	-37.115	88.723	-24.315	1.00	20.25
	ATOM	234	C	SER	30	-34.596	89.571	-23.121	1.00	31.61
40	ATOM	235	O	SER	30	-33.763	88.849	-23.659	1.00	35.14
	ATOM	236	N	PHE	31	-34.547	90.898	-23.175	1.00	32.12
	ATOM	237	CA	PHE	31	-33.474	91.593	-23.866	1.00	30.76
	ATOM	238	CB	PHE	31	-33.556	91.360	-25.379	1.00	27.55
	ATOM	239	CG	PHE	31	-34.426	92.342	-26.112	1.00	31.23
45	ATOM	240	CD1	PHE	31	-33.933	93.588	-26.482	1.00	32.00
	ATOM	241	CD2	PHE	31	-35.748	92.029	-26.421	1.00	27.89
	ATOM	242	CE1	PHE	31	-34.736	94.502	-27.145	1.00	29.70
	ATOM	243	CE2	PHE	31	-36.555	92.938	-27.085	1.00	26.15
	ATOM	244	CZ	PHE	31	-36.050	94.176	-27.447	1.00	28.30
50	ATOM	245	C	PHE	31	-33.539	93.079	-23.557	1.00	32.06
	ATOM	246	O	PHE	31	-34.606	93.630	-23.289	1.00	32.26
	ATOM	247	N	LYS	32	-32.383	93.719	-23.581	1.00	34.83
	ATOM	248	CA	LYS	32	-32.298	95.142	-23.325	1.00	37.13
	ATOM	249	CB	LYS	32	-31.622	95.395	-21.979	1.00	36.55
55	ATOM	250	CG	LYS	32	-31.437	96.854	-21.669	1.00	40.62
	ATOM	251	CD	LYS	32	-30.526	97.070	-20.486	1.00	44.56
	ATOM	252	CE	LYS	32	-30.136	98.548	-20.414	1.00	46.29
	ATOM	253	NZ	LYS	32	-29.333	98.885	-19.201	1.00	50.58
	ATOM	254	C	LYS	32	-31.477	95.747	-24.460	1.00	36.39
60	ATOM	255	O	LYS	32	-30.429	95.214	-24.828	1.00	38.69
	ATOM	256	N	ARG	33	-31.962	96.845	-25.028	1.00	37.90
	ATOM	257	CA	ARG	33	-31.261	97.497	-26.126	1.00	35.74
	ATOM	258	CB	ARG	33	-31.993	97.205	-27.431	1.00	36.79
	ATOM	259	CG	ARG	33	-31.428	97.881	-28.660	1.00	31.09
65	ATOM	260	CD	ARG	33	-32.073	97.292	-29.908	1.00	31.62
	ATOM	261	NE	ARG	33	-31.480	97.778	-31.151	1.00	38.16

	ATOM	262	CZ	ARG	33	-32.015	98.724	-31.914	1.00	39.15
	ATOM	263	NH1	ARG	33	-33.162	99.290	-31.565	1.00	43.41
	ATOM	264	NH2	ARG	33	-31.406	99.109	-33.023	1.00	40.65
	ATOM	265	C	ARG	33	-31.185	98.999	-25.880	1.00	37.71
5	ATOM	266	O	ARG	33	-32.212	99.674	-25.770	1.00	41.16
	ATOM	267	N	GLY	34	-29.971	99.529	-25.784	1.00	36.59
	ATOM	268	CA	GLY	34	-29.827	100.951	-25.552	1.00	36.72
	ATOM	269	C	GLY	34	-29.717	101.295	-24.081	1.00	40.01
	ATOM	270	O	GLY	34	-29.503	100.425	-23.238	1.00	39.85
10	ATOM	271	N	SER	35	-29.892	102.573	-23.774	1.00	38.74
	ATOM	272	CA	SER	35	-29.775	103.055	-22.406	1.00	40.23
	ATOM	273	CB	SER	35	-28.652	104.084	-22.349	1.00	43.02
	ATOM	274	OG	SER	35	-28.868	105.104	-23.323	1.00	48.05
	ATOM	275	C	SER	35	-31.037	103.678	-21.804	1.00	40.54
15	ATOM	276	O	SER	35	-31.127	103.816	-20.585	1.00	43.74
	ATOM	277	N	ALA	36	-32.000	104.047	-22.644	1.00	35.56
	ATOM	278	CA	ALA	36	-33.225	104.690	-22.181	1.00	29.62
	ATOM	279	CB	ALA	36	-34.072	105.077	-23.376	1.00	25.47
	ATOM	280	C	ALA	36	-34.072	103.904	-21.184	1.00	31.23
20	ATOM	281	O	ALA	36	-34.818	104.497	-20.414	1.00	29.56
	ATOM	282	N	LEU	37	-33.962	102.580	-21.188	1.00	34.09
	ATOM	283	CA	LEU	37	-34.760	101.753	-20.288	1.00	32.32
	ATOM	284	CB	LEU	37	-35.848	101.032	-21.080	1.00	26.47
	ATOM	285	CG	LEU	37	-36.822	101.949	-21.816	1.00	25.72
25	ATOM	286	CD1	LEU	37	-37.535	101.190	-22.907	1.00	24.61
	ATOM	287	CD2	LEU	37	-37.803	102.543	-20.833	1.00	27.53
	ATOM	288	C	LEU	37	-33.932	100.735	-19.523	1.00	35.16
	ATOM	289	O	LEU	37	-32.991	100.156	-20.053	1.00	36.47
	ATOM	290	N	GLU	38	-34.302	100.520	-18.270	1.00	38.30
30	ATOM	291	CA	GLU	38	-33.609	99.570	-17.414	1.00	41.17
	ATOM	292	CB	GLU	38	-32.698	100.304	-16.437	1.00	46.15
	ATOM	293	CG	GLU	38	-31.298	100.580	-16.932	1.00	50.01
	ATOM	294	CD	GLU	38	-30.560	101.549	-16.017	1.00	51.18
	ATOM	295	OE1	GLU	38	-30.722	101.426	-14.775	1.00	48.40
35	ATOM	296	OE2	GLU	38	-29.825	102.427	-16.541	1.00	50.23
	ATOM	297	C	GLU	38	-34.592	98.751	-16.600	1.00	42.69
	ATOM	298	O	GLU	38	-35.723	99.161	-16.372	1.00	42.55
	ATOM	299	N	GLU	39	-34.157	97.583	-16.156	1.00	44.75
	ATOM	300	CA	GLU	39	-35.018	96.751	-15.333	1.00	46.50
40	ATOM	301	CB	GLU	39	-34.769	95.273	-15.618	1.00	47.52
	ATOM	302	CG	GLU	39	-33.307	94.956	-15.846	1.00	58.50
	ATOM	303	CD	GLU	39	-33.051	93.468	-15.989	1.00	62.35
	ATOM	304	OE1	GLU	39	-31.914	93.090	-16.376	1.00	64.58
	ATOM	305	OE2	GLU	39	-33.983	92.679	-15.705	1.00	63.04
45	ATOM	306	C	GLU	39	-34.611	97.094	-13.912	1.00	44.56
	ATOM	307	O	GLU	39	-33.430	97.236	-13.623	1.00	48.11
	ATOM	308	N	LYS	40	-35.584	97.251	-13.029	1.00	40.71
	ATOM	309	CA	LYS	40	-35.288	97.593	-11.652	1.00	38.08
	ATOM	310	CB	LYS	40	-35.189	99.110	-11.496	1.00	41.28
50	ATOM	311	CG	LYS	40	-35.058	99.578	-10.052	1.00	41.52
	ATOM	312	CD	LYS	40	-35.299	101.077	-9.942	1.00	42.73
	ATOM	313	CE	LYS	40	-35.333	101.547	-8.498	1.00	41.97
	ATOM	314	NZ	LYS	40	-35.694	102.996	-8.419	1.00	45.94
	ATOM	315	C	LYS	40	-36.354	97.064	-10.716	1.00	39.66
55	ATOM	316	O	LYS	40	-37.501	97.499	-10.753	1.00	36.18
	ATOM	317	N	GLU	41	-35.959	96.115	-9.877	1.00	40.16
	ATOM	318	CA	GLU	41	-36.859	95.522	-8.900	1.00	42.80
	ATOM	319	CB	GLU	41	-37.063	96.508	-7.756	1.00	45.31
	ATOM	320	CG	GLU	41	-35.738	96.949	-7.145	1.00	56.07
60	ATOM	321	CD	GLU	41	-35.848	98.223	-6.313	1.00	59.42
	ATOM	322	OE1	GLU	41	-36.340	99.253	-6.837	1.00	63.71
	ATOM	323	OE2	GLU	41	-35.434	98.196	-5.133	1.00	63.71
	ATOM	324	C	GLU	41	-38.192	95.115	-9.508	1.00	38.93
	ATOM	325	O	GLU	41	-39.251	95.513	-9.032	1.00	39.03
65	ATOM	326	N	ASN	42	-38.117	94.333	-10.576	1.00	34.69
	ATOM	327	CA	ASN	42	-39.293	93.825	-11.262	1.00	32.96

	ATOM	328	CB	ASN	42	-40.190	93.083	-10.279	1.00	32.65
	ATOM	329	CG	ASN	42	-40.877	91.913	-10.911	1.00	35.50
	ATOM	330	OD1	ASN	42	-42.062	91.691	-10.704	1.00	40.26
	ATOM	331	ND2	ASN	42	-40.129	91.145	-11.689	1.00	29.28
5	ATOM	332	C	ASN	42	-40.124	94.868	-11.993	1.00	31.44
	ATOM	333	O	ASN	42	-41.279	94.623	-12.328	1.00	22.78
	ATOM	334	N	LYS	43	-39.533	96.025	-12.246	1.00	33.26
	ATOM	335	CA	LYS	43	-40.227	97.096	-12.941	1.00	33.68
	ATOM	336	CB	LYS	43	-40.611	98.201	-11.958	1.00	33.55
10	ATOM	337	CG	LYS	43	-41.656	97.787	-10.953	1.00	40.49
	ATOM	338	CD	LYS	43	-41.917	98.890	-9.959	1.00	45.39
	ATOM	339	CE	LYS	43	-40.689	99.150	-9.110	1.00	52.57
	ATOM	340	NZ	LYS	43	-40.966	100.160	-8.056	1.00	62.06
	ATOM	341	C	LYS	43	-39.330	97.670	-14.013	1.00	32.05
15	ATOM	342	O	LYS	43	-38.135	97.408	-14.029	1.00	37.32
	ATOM	343	N	ILE	44	-39.903	98.443	-14.923	1.00	31.31
	ATOM	344	CA	ILE	44	-39.096	99.047	-15.963	1.00	29.52
	ATOM	345	CB	ILE	44	-39.794	98.983	-17.322	1.00	26.38
	ATOM	346	CG2	ILE	44	-38.939	99.667	-18.369	1.00	26.09
20	ATOM	347	CG1	ILE	44	-40.010	97.521	-17.715	1.00	28.61
	ATOM	348	CD1	ILE	44	-40.667	97.327	-19.047	1.00	21.51
	ATOM	349	C	ILE	44	-38.822	100.492	-15.586	1.00	31.60
	ATOM	350	O	ILE	44	-39.743	101.277	-15.369	1.00	34.50
	ATOM	351	N	LEU	45	-37.543	100.831	-15.487	1.00	31.44
25	ATOM	352	CA	LEU	45	-37.125	102.178	-15.130	1.00	31.40
	ATOM	353	CB	LEU	45	-35.938	102.109	-14.173	1.00	33.60
	ATOM	354	CG	LEU	45	-35.305	103.445	-13.793	1.00	34.36
	ATOM	355	CD1	LEU	45	-36.274	104.269	-12.970	1.00	29.00
	ATOM	356	CD2	LEU	45	-34.030	103.180	-13.016	1.00	32.42
30	ATOM	357	C	LEU	45	-36.748	103.010	-16.352	1.00	27.80
	ATOM	358	O	LEU	45	-35.962	102.584	-17.191	1.00	27.94
	ATOM	359	N	VAL	46	-37.322	104.204	-16.440	1.00	28.00
	ATOM	360	CA	VAL	46	-37.049	105.115	-17.541	1.00	31.53
	ATOM	361	CB	VAL	46	-38.261	106.031	-17.780	1.00	27.89
35	ATOM	362	CG1	VAL	46	-38.008	106.946	-18.953	1.00	26.10
	ATOM	363	CG2	VAL	46	-39.486	105.193	-18.018	1.00	23.39
	ATOM	364	C	VAL	46	-35.817	105.971	-17.206	1.00	33.94
	ATOM	365	O	VAL	46	-35.783	106.638	-16.171	1.00	39.68
	ATOM	366	N	LYS	47	-34.813	105.953	-18.080	1.00	33.59
40	ATOM	367	CA	LYS	47	-33.594	106.724	-17.848	1.00	37.05
	ATOM	368	CB	LYS	47	-32.361	105.836	-18.044	1.00	34.49
	ATOM	369	CG	LYS	47	-32.440	104.508	-17.311	1.00	37.53
	ATOM	370	CD	LYS	47	-32.653	104.703	-15.816	1.00	41.69
	ATOM	371	CE	LYS	47	-31.336	104.680	-15.063	1.00	42.95
45	ATOM	372	NZ	LYS	47	-30.319	105.532	-15.720	1.00	44.50
	ATOM	373	C	LYS	47	-33.480	107.965	-18.735	1.00	37.34
	ATOM	374	O	LYS	47	-32.627	108.817	-18.508	1.00	43.04
	ATOM	375	N	GLU	48	-34.337	108.056	-19.741	1.00	36.99
	ATOM	376	CA	GLU	48	-34.356	109.196	-20.653	1.00	37.31
50	ATOM	377	CB	GLU	48	-33.745	108.832	-21.995	1.00	36.53
	ATOM	378	CG	GLU	48	-32.266	108.584	-21.983	1.00	46.74
	ATOM	379	CD	GLU	48	-31.760	108.182	-23.357	1.00	52.39
	ATOM	380	OE1	GLU	48	-32.235	108.780	-24.362	1.00	51.33
	ATOM	381	OE2	GLU	48	-30.890	107.276	-23.431	1.00	54.55
55	ATOM	382	C	GLU	48	-35.804	109.549	-20.887	1.00	36.94
	ATOM	383	O	GLU	48	-36.590	108.678	-21.236	1.00	40.21
	ATOM	384	N	THR	49	-36.177	110.812	-20.720	1.00	36.34
	ATOM	385	CA	THR	49	-37.577	111.147	-20.928	1.00	35.48
	ATOM	386	CB	THR	49	-37.966	112.480	-20.230	1.00	31.16
60	ATOM	387	OG1	THR	49	-37.924	113.548	-21.174	1.00	32.48
	ATOM	388	CG2	THR	49	-37.021	112.787	-19.095	1.00	30.36
	ATOM	389	C	THR	49	-37.897	111.214	-22.424	1.00	34.59
	ATOM	390	O	THR	49	-37.034	111.517	-23.246	1.00	35.13
	ATOM	391	N	GLY	50	-39.145	110.902	-22.764	1.00	32.49
65	ATOM	392	CA	GLY	50	-39.566	110.927	-24.148	1.00	28.17
	ATOM	393	C	GLY	50	-40.861	110.170	-24.332	1.00	29.99

	ATOM	394	O	GLY	50	-41.579	109.932	-23.374	1.00	29.76
	ATOM	395	N	TYR	51	-41.163	109.801	-25.572	1.00	31.68
	ATOM	396	CA	TYR	51	-42.373	109.048	-25.884	1.00	32.88
	ATOM	397	CB	TYR	51	-43.048	109.604	-27.143	1.00	35.97
5	ATOM	398	CG	TYR	51	-43.669	110.953	-26.925	1.00	43.03
	ATOM	399	CD1	TYR	51	-42.894	112.115	-26.961	1.00	44.15
	ATOM	400	CE1	TYR	51	-43.449	113.362	-26.667	1.00	47.81
	ATOM	401	CD2	TYR	51	-45.018	111.064	-26.598	1.00	47.08
	ATOM	402	CE2	TYR	51	-45.587	112.301	-26.298	1.00	51.79
10	ATOM	403	CZ	TYR	51	-44.798	113.448	-26.330	1.00	51.15
	ATOM	404	OH	TYR	51	-45.353	114.666	-25.982	1.00	52.20
	ATOM	405	C	TYR	51	-42.057	107.571	-26.079	1.00	31.97
	ATOM	406	O	TYR	51	-41.190	107.208	-26.871	1.00	30.01
	ATOM	407	N	PHE	52	-42.765	106.718	-25.349	1.00	31.50
15	ATOM	408	CA	PHE	52	-42.536	105.287	-25.445	1.00	30.41
	ATOM	409	CB	PHE	52	-42.029	104.735	-24.111	1.00	31.30
	ATOM	410	CG	PHE	52	-40.752	105.352	-23.633	1.00	29.70
	ATOM	411	CD1	PHE	52	-40.734	106.639	-23.124	1.00	26.20
	ATOM	412	CD2	PHE	52	-39.565	104.636	-23.683	1.00	28.38
20	ATOM	413	CE1	PHE	52	-39.557	107.201	-22.674	1.00	27.17
	ATOM	414	CE2	PHE	52	-38.381	105.192	-23.235	1.00	27.76
	ATOM	415	CZ	PHE	52	-38.376	106.475	-22.730	1.00	27.27
	ATOM	416	C	PHE	52	-43.770	104.498	-25.832	1.00	27.65
	ATOM	417	O	PHE	52	-44.887	104.858	-25.483	1.00	24.85
25	ATOM	418	N	PHE	53	-43.545	103.419	-26.565	1.00	27.18
	ATOM	419	CA	PHE	53	-44.608	102.509	-26.959	1.00	27.05
	ATOM	420	CB	PHE	53	-44.338	101.917	-28.336	1.00	28.76
	ATOM	421	CG	PHE	53	-45.289	100.830	-28.719	1.00	28.00
	ATOM	422	CD1	PHE	53	-46.612	101.117	-29.022	1.00	26.49
30	ATOM	423	CD2	PHE	53	-44.869	99.508	-28.761	1.00	28.04
	ATOM	424	CE1	PHE	53	-47.500	100.103	-29.363	1.00	26.54
	ATOM	425	CE2	PHE	53	-45.756	98.490	-29.100	1.00	25.78
	ATOM	426	CZ	PHE	53	-47.071	98.792	-29.400	1.00	24.39
	ATOM	427	C	PHE	53	-44.499	101.413	-25.902	1.00	27.56
35	ATOM	428	O	PHE	53	-43.452	100.789	-25.765	1.00	25.39
	ATOM	429	N	ILE	54	-45.572	101.197	-25.147	1.00	28.87
	ATOM	430	CA	ILE	54	-45.572	100.204	-24.079	1.00	28.95
	ATOM	431	CB	ILE	54	-45.921	100.881	-22.738	1.00	29.60
	ATOM	432	CG2	ILE	54	-45.651	99.925	-21.587	1.00	27.42
40	ATOM	433	CG1	ILE	54	-45.078	102.148	-22.573	1.00	26.41
	ATOM	434	CD1	ILE	54	-45.542	103.069	-21.487	1.00	28.70
	ATOM	435	C	ILE	54	-46.564	99.076	-24.363	1.00	28.57
	ATOM	436	O	ILE	54	-47.684	99.316	-24.801	1.00	29.17
	ATOM	437	N	TYR	55	-46.156	97.839	-24.109	1.00	26.97
45	ATOM	438	CA	TYR	55	-47.031	96.706	-24.382	1.00	24.22
	ATOM	439	CB	TYR	55	-46.698	96.123	-25.756	1.00	21.82
	ATOM	440	CG	TYR	55	-45.271	95.658	-25.904	1.00	27.68
	ATOM	441	CD1	TYR	55	-44.907	94.350	-25.599	1.00	27.01
	ATOM	442	CE1	TYR	55	-43.594	93.931	-25.714	1.00	27.44
50	ATOM	443	CD2	TYR	55	-44.277	96.533	-26.331	1.00	27.16
	ATOM	444	CE2	TYR	55	-42.959	96.121	-26.446	1.00	25.39
	ATOM	445	CZ	TYR	55	-42.625	94.823	-26.136	1.00	29.02
	ATOM	446	OH	TYR	55	-41.318	94.421	-26.240	1.00	31.82
	ATOM	447	C	TYR	55	-46.953	95.624	-23.327	1.00	25.16
55	ATOM	448	O	TYR	55	-46.031	95.586	-22.544	1.00	28.47
	ATOM	449	N	GLY	56	-47.940	94.745	-23.301	1.00	27.44
	ATOM	450	CA	GLY	56	-47.940	93.680	-22.322	1.00	22.07
	ATOM	451	C	GLY	56	-48.995	92.635	-22.611	1.00	25.83
	ATOM	452	O	GLY	56	-50.062	92.950	-23.127	1.00	26.00
60	ATOM	453	N	GLN	57	-48.682	91.384	-22.297	1.00	22.56
	ATOM	454	CA	GLN	57	-49.608	90.280	-22.493	1.00	20.91
	ATOM	455	CB	GLN	57	-49.340	89.540	-23.804	1.00	17.87
	ATOM	456	CG	GLN	57	-50.211	88.300	-23.965	1.00	16.86
	ATOM	457	CD	GLN	57	-50.028	87.594	-25.292	1.00	22.06
65	ATOM	458	OE1	GLN	57	-50.333	88.144	-26.345	1.00	27.26
	ATOM	459	NE2	GLN	57	-49.533	86.364	-25.246	1.00	20.20

	ATOM	460	C	GLN	57	-49.465	89.309	-21.340	1.00	24.23
	ATOM	461	O	GLN	57	-48.366	89.107	-20.827	1.00	25.16
	ATOM	462	N	VAL	58	-50.587	88.716	-20.942	1.00	24.62
	ATOM	463	CA	VAL	58	-50.634	87.748	-19.853	1.00	26.68
5	ATOM	464	CB	VAL	58	-51.155	88.393	-18.545	1.00	25.55
	ATOM	465	CG1	VAL	58	-51.391	87.332	-17.499	1.00	15.69
	ATOM	466	CG2	VAL	58	-50.166	89.417	-18.030	1.00	25.77
	ATOM	467	C	VAL	58	-51.607	86.656	-20.249	1.00	31.06
	ATOM	468	O	VAL	58	-52.638	86.952	-20.837	1.00	27.74
10	ATOM	469	N	LEU	59	-51.282	85.401	-19.936	1.00	33.82
	ATOM	470	CA	LEU	59	-52.176	84.276	-20.232	1.00	29.85
	ATOM	471	CB	LEU	59	-51.396	83.052	-20.702	1.00	29.62
	ATOM	472	CG	LEU	59	-52.150	82.016	-21.546	1.00	28.11
	ATOM	473	CD1	LEU	59	-51.337	80.740	-21.611	1.00	26.67
15	ATOM	474	CD2	LEU	59	-53.507	81.725	-20.977	1.00	22.45
	ATOM	475	C	LEU	59	-52.918	83.920	-18.946	1.00	30.98
	ATOM	476	O	LEU	59	-52.312	83.480	-17.972	1.00	32.52
	ATOM	477	N	TYR	60	-54.232	84.117	-18.946	1.00	33.01
	ATOM	478	CA	TYR	60	-55.038	83.814	-17.772	1.00	34.51
20	ATOM	479	CB	TYR	60	-56.180	84.818	-17.648	1.00	40.10
	ATOM	480	CG	TYR	60	-55.691	86.222	-17.464	1.00	45.37
	ATOM	481	CD1	TYR	60	-55.759	87.150	-18.504	1.00	47.13
	ATOM	482	CE1	TYR	60	-55.226	88.423	-18.356	1.00	50.28
	ATOM	483	CD2	TYR	60	-55.087	86.604	-16.273	1.00	44.35
25	ATOM	484	CE2	TYR	60	-54.552	87.867	-16.111	1.00	48.81
	ATOM	485	CZ	TYR	60	-54.619	88.769	-17.153	1.00	51.91
	ATOM	486	OH	TYR	60	-54.046	90.006	-16.996	1.00	53.77
	ATOM	487	C	TYR	60	-55.603	82.404	-17.780	1.00	35.49
	ATOM	488	O	TYR	60	-56.307	82.004	-18.701	1.00	36.13
30	ATOM	489	N	THR	61	-55.279	81.653	-16.737	1.00	36.19
	ATOM	490	CA	THR	61	-55.757	80.291	-16.593	1.00	35.36
	ATOM	491	CB	THR	61	-54.589	79.305	-16.560	1.00	34.22
	ATOM	492	OG1	THR	61	-53.652	79.710	-15.558	1.00	34.22
	ATOM	493	CG2	THR	61	-53.898	79.255	-17.909	1.00	25.45
35	ATOM	494	C	THR	61	-56.528	80.233	-15.282	1.00	37.82
	ATOM	495	O	THR	61	-56.544	79.221	-14.589	1.00	43.70
	ATOM	496	N	ASP	62	-57.163	81.352	-14.959	1.00	37.64
	ATOM	497	CA	ASP	62	-57.948	81.520	-13.750	1.00	38.08
	ATOM	498	CB	ASP	62	-57.521	82.829	-13.092	1.00	38.42
40	ATOM	499	CG	ASP	62	-58.129	83.031	-11.734	1.00	41.25
	ATOM	500	OD1	ASP	62	-57.411	83.540	-10.841	1.00	38.28
	ATOM	501	OD2	ASP	62	-59.320	82.694	-11.571	1.00	41.37
	ATOM	502	C	ASP	62	-59.415	81.572	-14.167	1.00	40.90
	ATOM	503	O	ASP	62	-59.732	82.157	-15.194	1.00	46.73
45	ATOM	504	N	LYS	63	-60.316	80.971	-13.395	1.00	42.62
	ATOM	505	CA	LYS	63	-61.726	80.987	-13.785	1.00	40.92
	ATOM	506	CB	LYS	63	-62.353	79.609	-13.566	1.00	42.79
	ATOM	507	CG	LYS	63	-62.294	79.098	-12.143	1.00	45.91
	ATOM	508	CD	LYS	63	-62.746	77.639	-12.083	1.00	46.22
50	ATOM	509	CE	LYS	63	-61.817	76.737	-12.918	1.00	49.97
	ATOM	510	NZ	LYS	63	-62.323	75.331	-13.090	1.00	49.98
	ATOM	511	C	LYS	63	-62.583	82.051	-13.107	1.00	42.59
	ATOM	512	O	LYS	63	-63.803	81.947	-13.091	1.00	42.46
	ATOM	513	N	THR	64	-61.941	83.090	-12.589	1.00	44.06
55	ATOM	514	CA	THR	64	-62.631	84.174	-11.905	1.00	44.94
	ATOM	515	CB	THR	64	-61.612	85.070	-11.159	1.00	43.79
	ATOM	516	OG1	THR	64	-60.824	84.259	-10.282	1.00	46.20
	ATOM	517	CG2	THR	64	-62.319	86.133	-10.325	1.00	42.80
	ATOM	518	C	THR	64	-63.535	85.064	-12.774	1.00	45.82
60	ATOM	519	O	THR	64	-63.514	86.275	-12.641	1.00	53.10
	ATOM	520	N	TYR	65	-64.333	84.473	-13.649	1.00	45.11
	ATOM	521	CA	TYR	65	-65.271	85.212	-14.507	1.00	45.72
	ATOM	522	CB	TYR	65	-66.492	85.665	-13.685	1.00	40.46
	ATOM	523	CG	TYR	65	-66.423	87.065	-13.116	1.00	41.90
65	ATOM	524	CD1	TYR	65	-66.849	88.168	-13.859	1.00	42.47
	ATOM	525	CE1	TYR	65	-66.777	89.472	-13.338	1.00	41.83

	ATOM	526	CD2	TYR	65	-65.923	87.295	-11.830	1.00	41.65
	ATOM	527	CE2	TYR	65	-65.844	88.593	-11.297	1.00	40.28
	ATOM	528	CZ	TYR	65	-66.272	89.672	-12.059	1.00	41.90
	ATOM	529	OH	TYR	65	-66.197	90.942	-11.543	1.00	43.84
5	ATOM	530	C	TYR	65	-64.725	86.400	-15.315	1.00	45.31
	ATOM	531	O	TYR	65	-65.224	86.698	-16.409	1.00	49.74
	ATOM	532	N	ALA	66	-63.718	87.083	-14.787	1.00	42.32
	ATOM	533	CA	ALA	66	-63.128	88.226	-15.470	1.00	39.57
	ATOM	534	CB	ALA	66	-63.973	89.462	-15.250	1.00	35.36
10	ATOM	535	C	ALA	66	-61.719	88.460	-14.957	1.00	39.85
	ATOM	536	O	ALA	66	-61.507	88.664	-13.760	1.00	37.60
	ATOM	537	N	MET	67	-60.753	88.410	-15.865	1.00	41.88
	ATOM	538	CA	MET	67	-59.359	88.622	-15.506	1.00	41.48
	ATOM	539	CB	MET	67	-58.537	87.346	-15.721	1.00	39.98
15	ATOM	540	CG	MET	67	-58.862	86.206	-14.764	1.00	37.31
	ATOM	541	SD	MET	67	-58.615	86.642	-13.037	1.00	35.46
	ATOM	542	CE	MET	67	-56.841	86.441	-12.860	1.00	34.06
	ATOM	543	C	MET	67	-58.792	89.749	-16.352	1.00	39.50
	ATOM	544	O	MET	67	-59.395	90.159	-17.346	1.00	38.44
20	ATOM	545	N	GLY	68	-57.630	90.248	-15.950	1.00	39.84
	ATOM	546	CA	GLY	68	-57.000	91.325	-16.687	1.00	37.61
	ATOM	547	C	GLY	68	-55.821	91.907	-15.945	1.00	33.76
	ATOM	548	O	GLY	68	-55.574	91.557	-14.796	1.00	35.58
	ATOM	549	N	HIS	69	-55.081	92.787	-16.605	1.00	35.07
25	ATOM	550	CA	HIS	69	-53.929	93.417	-15.982	1.00	33.70
	ATOM	551	CB	HIS	69	-52.619	92.753	-16.447	1.00	37.02
	ATOM	552	CG	HIS	69	-52.453	92.680	-17.933	1.00	34.65
	ATOM	553	CD2	HIS	69	-51.665	93.386	-18.775	1.00	35.63
	ATOM	554	ND1	HIS	69	-53.140	91.781	-18.719	1.00	37.68
30	ATOM	555	CE1	HIS	69	-52.782	91.933	-19.981	1.00	32.64
	ATOM	556	NE2	HIS	69	-51.886	92.901	-20.042	1.00	35.90
	ATOM	557	C	HIS	69	-53.887	94.912	-16.268	1.00	33.98
	ATOM	558	O	HIS	69	-54.635	95.414	-17.092	1.00	30.94
	ATOM	559	N	LEU	70	-53.006	95.618	-15.573	1.00	35.35
35	ATOM	560	CA	LEU	70	-52.862	97.056	-15.749	1.00	30.76
	ATOM	561	CB	LEU	70	-53.275	97.789	-14.478	1.00	29.59
	ATOM	562	CG	LEU	70	-54.491	97.298	-13.699	1.00	30.71
	ATOM	563	CD1	LEU	70	-54.543	98.004	-12.366	1.00	25.60
	ATOM	564	CD2	LEU	70	-55.750	97.542	-14.497	1.00	27.39
40	ATOM	565	C	LEU	70	-51.404	97.409	-16.023	1.00	33.19
	ATOM	566	O	LEU	70	-50.497	96.855	-15.401	1.00	37.07
	ATOM	567	N	ILE	71	-51.169	98.310	-16.970	1.00	30.93
	ATOM	568	CA	ILE	71	-49.810	98.760	-17.227	1.00	27.94
	ATOM	569	CB	ILE	71	-49.530	98.948	-18.720	1.00	28.22
45	ATOM	570	CG2	ILE	71	-48.249	99.731	-18.916	1.00	24.54
	ATOM	571	CG1	ILE	71	-49.399	97.584	-19.388	1.00	26.34
	ATOM	572	CD1	ILE	71	-49.319	97.653	-20.888	1.00	37.50
	ATOM	573	C	ILE	71	-49.825	100.105	-16.520	1.00	30.32
	ATOM	574	O	ILE	71	-50.567	100.998	-16.912	1.00	23.07
50	ATOM	575	N	GLN	72	-49.033	100.240	-15.460	1.00	29.74
	ATOM	576	CA	GLN	72	-49.033	101.483	-14.702	1.00	27.85
	ATOM	577	CB	GLN	72	-49.392	101.184	-13.255	1.00	26.70
	ATOM	578	CG	GLN	72	-50.631	100.344	-13.132	1.00	28.28
	ATOM	579	CD	GLN	72	-51.097	100.216	-11.710	1.00	33.11
55	ATOM	580	OE1	GLN	72	-50.333	99.822	-10.834	1.00	41.22
	ATOM	581	NE2	GLN	72	-52.359	100.543	-11.466	1.00	30.18
	ATOM	582	C	GLN	72	-47.748	102.283	-14.746	1.00	29.85
	ATOM	583	O	GLN	72	-46.671	101.750	-15.019	1.00	28.20
	ATOM	584	N	ARG	73	-47.881	103.572	-14.456	1.00	30.90
60	ATOM	585	CA	ARG	73	-46.753	104.487	-14.444	1.00	30.79
	ATOM	586	CB	ARG	73	-46.940	105.534	-15.534	1.00	29.18
	ATOM	587	CG	ARG	73	-45.874	106.587	-15.556	1.00	28.66
	ATOM	588	CD	ARG	73	-46.389	107.829	-16.221	1.00	29.25
	ATOM	589	NE	ARG	73	-45.363	108.857	-16.306	1.00	35.78
65	ATOM	590	CZ	ARG	73	-45.613	110.140	-16.515	1.00	35.27
	ATOM	591	NH1	ARG	73	-46.862	110.561	-16.654	1.00	34.95

	ATOM	592	NH2	ARG	73	-44.612	110.999	-16.589	1.00	33.33
	ATOM	593	C	ARG	73	-46.605	105.191	-13.097	1.00	32.42
	ATOM	594	O	ARG	73	-47.563	105.760	-12.586	1.00	34.19
	ATOM	595	N	LYS	74	-45.408	105.132	-12.522	1.00	35.90
5	ATOM	596	CA	LYS	74	-45.126	105.811	-11.257	1.00	38.57
	ATOM	597	CB	LYS	74	-44.259	104.945	-10.338	1.00	43.52
	ATOM	598	CG	LYS	74	-44.980	103.766	-9.707	1.00	52.89
	ATOM	599	CD	LYS	74	-44.039	102.907	-8.830	1.00	58.14
	ATOM	600	CE	LYS	74	-44.789	101.721	-8.222	1.00	56.75
10	ATOM	601	NZ	LYS	74	-43.897	100.847	-7.404	1.00	63.98
	ATOM	602	C	LYS	74	-44.353	107.081	-11.595	1.00	40.11
	ATOM	603	O	LYS	74	-43.168	107.017	-11.944	1.00	37.55
	ATOM	604	N	LYS	75	-45.019	108.226	-11.491	1.00	34.48
	ATOM	605	CA	LYS	75	-44.393	109.506	-11.798	1.00	34.24
15	ATOM	606	CB	LYS	75	-45.446	110.617	-11.792	1.00	36.16
	ATOM	607	CG	LYS	75	-46.631	110.439	-12.739	1.00	35.99
	ATOM	608	CD	LYS	75	-47.588	111.629	-12.590	1.00	40.91
	ATOM	609	CE	LYS	75	-48.782	111.527	-13.530	1.00	46.16
	ATOM	610	NZ	LYS	75	-49.786	112.638	-13.390	1.00	46.03
20	ATOM	611	C	LYS	75	-43.303	109.859	-10.780	1.00	30.92
	ATOM	612	O	LYS	75	-43.474	109.632	-9.597	1.00	27.47
	ATOM	613	N	VAL	76	-42.190	110.417	-11.250	1.00	32.82
	ATOM	614	CA	VAL	76	-41.096	110.830	-10.367	1.00	31.51
	ATOM	615	CB	VAL	76	-39.780	111.095	-11.120	1.00	29.19
25	ATOM	616	CG1	VAL	76	-38.615	110.688	-10.264	1.00	27.09
	ATOM	617	CG2	VAL	76	-39.768	110.386	-12.428	1.00	34.94
	ATOM	618	C	VAL	76	-41.480	112.164	-9.758	1.00	33.40
	ATOM	619	O	VAL	76	-41.238	112.419	-8.584	1.00	32.92
	ATOM	620	N	HIS	77	-42.066	113.018	-10.592	1.00	34.88
30	ATOM	621	CA	HIS	77	-42.495	114.343	-10.183	1.00	34.59
	ATOM	622	CB	HIS	77	-42.022	115.385	-11.201	1.00	35.92
	ATOM	623	CG	HIS	77	-40.537	115.406	-11.401	1.00	34.17
	ATOM	624	CD2	HIS	77	-39.786	115.806	-12.452	1.00	31.44
	ATOM	625	ND1	HIS	77	-39.647	115.005	-10.426	1.00	35.05
35	ATOM	626	CE1	HIS	77	-38.412	115.154	-10.866	1.00	29.84
	ATOM	627	NE2	HIS	77	-38.469	115.640	-12.091	1.00	38.35
	ATOM	628	C	HIS	77	-44.010	114.362	-10.066	1.00	35.63
	ATOM	629	O	HIS	77	-44.706	113.792	-10.907	1.00	40.24
	ATOM	630	N	VAL	78	-44.523	115.038	-9.040	1.00	37.37
40	ATOM	631	CA	VAL	78	-45.958	115.059	-8.822	1.00	41.32
	ATOM	632	CB	VAL	78	-46.290	114.374	-7.482	1.00	43.10
	ATOM	633	CG1	VAL	78	-47.786	114.263	-7.314	1.00	45.01
	ATOM	634	CG2	VAL	78	-45.688	112.981	-7.454	1.00	44.98
	ATOM	635	C	VAL	78	-46.740	116.374	-8.935	1.00	43.02
45	ATOM	636	O	VAL	78	-47.557	116.508	-9.845	1.00	49.79
	ATOM	637	N	PHE	79	-46.538	117.337	-8.046	1.00	41.33
	ATOM	638	CA	PHE	79	-47.306	118.603	-8.137	1.00	45.00
	ATOM	639	CB	PHE	79	-47.445	119.118	-9.584	1.00	40.51
	ATOM	640	CG	PHE	79	-46.151	119.295	-10.305	1.00	39.52
50	ATOM	641	CD1	PHE	79	-45.763	118.385	-11.282	1.00	36.80
	ATOM	642	CD2	PHE	79	-45.323	120.371	-10.015	1.00	37.89
	ATOM	643	CE1	PHE	79	-44.567	118.544	-11.962	1.00	37.52
	ATOM	644	CE2	PHE	79	-44.125	120.538	-10.692	1.00	35.59
	ATOM	645	CZ	PHE	79	-43.745	119.623	-11.668	1.00	37.58
55	ATOM	646	C	PHE	79	-48.735	118.519	-7.601	1.00	43.50
	ATOM	647	O	PHE	79	-49.536	117.697	-8.049	1.00	37.85
	ATOM	648	N	GLY	80	-49.051	119.409	-6.667	1.00	46.65
	ATOM	649	CA	GLY	80	-50.382	119.475	-6.095	1.00	48.38
	ATOM	650	C	GLY	80	-51.027	118.174	-5.672	1.00	47.89
60	ATOM	651	O	GLY	80	-50.414	117.356	-4.980	1.00	51.15
	ATOM	652	N	ASP	81	-52.276	117.983	-6.093	1.00	45.07
	ATOM	653	CA	ASP	81	-53.028	116.795	-5.733	1.00	43.99
	ATOM	654	CB	ASP	81	-54.482	117.170	-5.416	1.00	45.66
	ATOM	655	CG	ASP	81	-55.261	117.614	-6.645	1.00	47.49
65	ATOM	656	OD1	ASP	81	-54.643	117.893	-7.696	1.00	51.90
	ATOM	657	OD2	ASP	81	-56.501	117.695	-6.557	1.00	48.08

	ATOM	658	C	ASP	81	-52.991	115.694	-6.778	1.00	42.29
	ATOM	659	O	ASP	81	-53.884	114.848	-6.819	1.00	39.94
	ATOM	660	N	GLU	82	-51.964	115.701	-7.621	1.00	39.91
	ATOM	661	CA	GLU	82	-51.820	114.662	-8.639	1.00	36.58
5	ATOM	662	CB	GLU	82	-50.601	114.904	-9.529	1.00	34.91
	ATOM	663	CG	GLU	82	-50.733	115.919	-10.619	1.00	37.82
	ATOM	664	CD	GLU	82	-49.752	115.649	-11.748	1.00	40.75
	ATOM	665	OE1	GLU	82	-48.578	115.339	-11.462	1.00	38.15
	ATOM	666	OE2	GLU	82	-50.151	115.739	-12.928	1.00	47.99
10	ATOM	667	C	GLU	82	-51.588	113.321	-7.956	1.00	36.58
	ATOM	668	O	GLU	82	-50.998	113.259	-6.881	1.00	38.40
	ATOM	669	N	LEU	83	-52.054	112.251	-8.584	1.00	33.83
	ATOM	670	CA	LEU	83	-51.822	110.910	-8.072	1.00	32.70
	ATOM	671	CB	LEU	83	-52.924	109.955	-8.529	1.00	30.78
15	ATOM	672	CG	LEU	83	-54.122	109.749	-7.604	1.00	29.26
	ATOM	673	CD1	LEU	83	-54.587	111.061	-7.025	1.00	29.60
	ATOM	674	CD2	LEU	83	-55.230	109.084	-8.380	1.00	28.65
	ATOM	675	C	LEU	83	-50.517	110.549	-8.749	1.00	33.88
	ATOM	676	O	LEU	83	-50.371	110.763	-9.947	1.00	38.46
20	ATOM	677	N	SER	84	-49.561	110.021	-7.998	1.00	34.94
	ATOM	678	CA	SER	84	-48.278	109.678	-8.592	1.00	39.77
	ATOM	679	CB	SER	84	-47.185	109.659	-7.525	1.00	38.92
	ATOM	680	OG	SER	84	-47.498	108.732	-6.515	1.00	43.11
	ATOM	681	C	SER	84	-48.295	108.350	-9.343	1.00	39.89
25	ATOM	682	O	SER	84	-47.359	108.020	-10.069	1.00	42.52
	ATOM	683	N	LEU	85	-49.369	107.593	-9.171	1.00	40.92
	ATOM	684	CA	LEU	85	-49.510	106.316	-9.843	1.00	34.66
	ATOM	685	CB	LEU	85	-49.843	105.225	-8.825	1.00	35.12
	ATOM	686	CG	LEU	85	-49.784	103.736	-9.195	1.00	36.92
30	ATOM	687	CD1	LEU	85	-50.879	103.396	-10.177	1.00	38.24
	ATOM	688	CD2	LEU	85	-48.415	103.401	-9.767	1.00	36.08
	ATOM	689	C	LEU	85	-50.632	106.464	-10.853	1.00	35.30
	ATOM	690	O	LEU	85	-51.784	106.694	-10.496	1.00	32.85
	ATOM	691	N	VAL	86	-50.283	106.357	-12.126	1.00	33.26
35	ATOM	692	CA	VAL	86	-51.261	106.474	-13.195	1.00	34.57
	ATOM	693	CB	VAL	86	-50.856	107.558	-14.235	1.00	35.39
	ATOM	694	CG1	VAL	86	-51.804	107.533	-15.418	1.00	31.44
	ATOM	695	CG2	VAL	86	-50.872	108.930	-13.591	1.00	38.08
	ATOM	696	C	VAL	86	-51.358	105.154	-13.922	1.00	32.85
40	ATOM	697	O	VAL	86	-50.342	104.548	-14.252	1.00	35.10
	ATOM	698	N	THR	87	-52.571	104.690	-14.164	1.00	33.41
	ATOM	699	CA	THR	87	-52.705	103.457	-14.903	1.00	33.56
	ATOM	700	CB	THR	87	-53.715	102.483	-14.221	1.00	32.70
	ATOM	701	OG1	THR	87	-54.523	101.836	-15.212	1.00	34.29
45	ATOM	702	CG2	THR	87	-54.571	103.212	-13.219	1.00	34.95
	ATOM	703	C	THR	87	-53.080	103.755	-16.360	1.00	30.57
	ATOM	704	O	THR	87	-54.133	104.320	-16.648	1.00	28.08
	ATOM	705	N	LEU	88	-52.158	103.424	-17.257	1.00	28.42
	ATOM	706	CA	LEU	88	-52.348	103.588	-18.686	1.00	31.31
50	ATOM	707	CB	LEU	88	-51.011	103.836	-19.398	1.00	28.21
	ATOM	708	CG	LEU	88	-49.928	104.880	-19.105	1.00	29.12
	ATOM	709	CD1	LEU	88	-50.380	105.865	-18.086	1.00	31.12
	ATOM	710	CD2	LEU	88	-48.674	104.173	-18.650	1.00	28.72
	ATOM	711	C	LEU	88	-52.883	102.238	-19.192	1.00	41.15
55	ATOM	712	O	LEU	88	-52.554	101.179	-18.650	1.00	52.30
	ATOM	713	N	PHE	89	-53.723	102.247	-20.210	1.00	39.23
	ATOM	714	CA	PHE	89	-54.178	100.972	-20.771	1.00	42.27
	ATOM	715	CB	PHE	89	-52.951	100.264	-21.352	1.00	35.63
	ATOM	716	CG	PHE	89	-52.031	101.211	-22.078	1.00	35.06
60	ATOM	717	CD1	PHE	89	-50.653	101.142	-21.901	1.00	31.56
	ATOM	718	CD2	PHE	89	-52.563	102.261	-22.848	1.00	30.96
	ATOM	719	CE1	PHE	89	-49.817	102.112	-22.464	1.00	33.76
	ATOM	720	CE2	PHE	89	-51.741	103.231	-23.410	1.00	27.38
	ATOM	721	CZ	PHE	89	-50.367	103.161	-23.216	1.00	30.29
65	ATOM	722	C	PHE	89	-55.029	100.023	-19.898	1.00	39.99
	ATOM	723	O	PHE	89	-56.144	100.384	-19.508	1.00	45.34

	ATOM	724	N	ARG	90	-54.559	98.821	-19.601	1.00	38.92
	ATOM	725	CA	ARG	90	-55.399	97.892	-18.822	1.00	41.66
	ATOM	726	CB	ARG	90	-56.117	98.627	-17.672	1.00	39.24
	ATOM	727	CG	ARG	90	-57.549	98.137	-17.380	1.00	29.41
5	ATOM	728	CD	ARG	90	-58.199	98.891	-16.204	1.00	31.13
	ATOM	729	NE	ARG	90	-59.637	98.628	-16.102	1.00	31.75
	ATOM	730	CZ	ARG	90	-60.598	99.534	-16.303	1.00	28.86
	ATOM	731	NH1	ARG	90	-60.302	100.786	-16.612	1.00	25.48
	ATOM	732	NH2	ARG	90	-61.869	99.177	-16.225	1.00	29.84
10	ATOM	733	C	ARG	90	-56.472	97.158	-19.650	1.00	39.23
	ATOM	734	O	ARG	90	-57.235	97.782	-20.386	1.00	31.33
	ATOM	735	N	CYS	91	-56.524	95.830	-19.524	1.00	40.12
	ATOM	736	CA	CYS	91	-57.534	95.053	-20.225	1.00	37.09
	ATOM	737	C	CYS	91	-58.222	93.978	-19.434	1.00	37.21
15	ATOM	738	O	CYS	91	-57.787	93.612	-18.349	1.00	37.46
	ATOM	739	CB	CYS	91	-57.003	94.438	-21.499	1.00	40.13
	ATOM	740	SG	CYS	91	-55.630	93.231	-21.516	1.00	41.64
	ATOM	741	N	ILE	92	-59.309	93.472	-20.006	1.00	34.02
	ATOM	742	CA	ILE	92	-60.122	92.452	-19.360	1.00	33.22
20	ATOM	743	CB	ILE	92	-61.411	93.083	-18.793	1.00	32.88
	ATOM	744	CG2	ILE	92	-62.163	92.071	-17.955	1.00	32.61
	ATOM	745	CG1	ILE	92	-61.054	94.301	-17.937	1.00	34.64
	ATOM	746	CD1	ILE	92	-62.250	95.122	-17.483	1.00	32.30
	ATOM	747	C	ILE	92	-60.520	91.323	-20.303	1.00	30.78
25	ATOM	748	O	ILE	92	-60.602	91.513	-21.498	1.00	33.43
	ATOM	749	N	GLN	93	-60.740	90.137	-19.749	1.00	29.35
	ATOM	750	CA	GLN	93	-61.167	88.981	-20.521	1.00	28.69
	ATOM	751	CB	GLN	93	-59.994	88.068	-20.879	1.00	28.17
	ATOM	752	CG	GLN	93	-59.180	88.427	-22.125	1.00	25.87
30	ATOM	753	CD	GLN	93	-59.988	88.431	-23.405	1.00	24.22
	ATOM	754	OE1	GLN	93	-60.565	89.440	-23.775	1.00	28.90
	ATOM	755	NE2	GLN	93	-60.031	87.294	-24.086	1.00	25.29
	ATOM	756	C	GLN	93	-62.121	88.192	-19.654	1.00	31.44
	ATOM	757	O	GLN	93	-61.791	87.860	-18.525	1.00	32.07
35	ATOM	758	N	ASN	94	-63.317	87.916	-20.155	1.00	35.77
	ATOM	759	CA	ASN	94	-64.253	87.112	-19.392	1.00	33.56
	ATOM	760	CB	ASN	94	-65.640	87.124	-20.036	1.00	34.70
	ATOM	761	CG	ASN	94	-66.438	88.352	-19.679	1.00	33.01
	ATOM	762	OD1	ASN	94	-66.583	88.686	-18.515	1.00	30.56
40	ATOM	763	ND2	ASN	94	-66.976	89.020	-20.680	1.00	36.61
	ATOM	764	C	ASN	94	-63.672	85.698	-19.415	1.00	35.80
	ATOM	765	O	ASN	94	-63.096	85.271	-20.418	1.00	36.63
	ATOM	766	N	MET	95	-63.808	84.975	-18.311	1.00	35.73
	ATOM	767	CA	MET	95	-63.273	83.621	-18.234	1.00	34.73
45	ATOM	768	CB	MET	95	-62.381	83.491	-16.995	1.00	32.97
	ATOM	769	CG	MET	95	-61.267	84.523	-16.893	1.00	27.35
	ATOM	770	SD	MET	95	-60.193	84.589	-18.341	1.00	25.14
	ATOM	771	CE	MET	95	-59.258	83.104	-18.201	1.00	15.55
	ATOM	772	C	MET	95	-64.372	82.558	-18.187	1.00	36.36
50	ATOM	773	O	MET	95	-65.486	82.826	-17.727	1.00	40.70
	ATOM	774	N	PRO	96	-64.082	81.347	-18.691	1.00	36.06
	ATOM	775	CD	PRO	96	-62.962	81.047	-19.584	1.00	35.32
	ATOM	776	CA	PRO	96	-65.032	80.235	-18.698	1.00	40.00
	ATOM	777	CB	PRO	96	-64.553	79.360	-19.852	1.00	33.97
55	ATOM	778	CG	PRO	96	-63.649	80.232	-20.626	1.00	37.05
	ATOM	779	C	PRO	96	-64.877	79.506	-17.365	1.00	46.34
	ATOM	780	O	PRO	96	-63.917	79.748	-16.635	1.00	47.59
	ATOM	781	N	GLU	97	-65.802	78.605	-17.050	1.00	52.14
	ATOM	782	CA	GLU	97	-65.728	77.866	-15.796	1.00	57.27
60	ATOM	783	CB	GLU	97	-67.122	77.416	-15.358	1.00	62.29
	ATOM	784	CG	GLU	97	-67.261	77.146	-13.863	1.00	72.31
	ATOM	785	CD	GLU	97	-67.908	78.322	-13.118	1.00	79.34
	ATOM	786	OE1	GLU	97	-69.093	78.658	-13.428	1.00	84.94
	ATOM	787	OE2	GLU	97	-67.232	78.912	-12.228	1.00	81.39
65	ATOM	788	C	GLU	97	-64.847	76.637	-15.993	1.00	56.67
	ATOM	789	O	GLU	97	-64.099	76.241	-15.095	1.00	59.18

	ATOM	790	N	THR	98	-64.925	76.052	-17.182	1.00	57.55
	ATOM	791	CA	THR	98	-64.176	74.842	-17.503	1.00	59.71
	ATOM	792	CB	THR	98	-64.663	74.248	-18.819	1.00	59.24
	ATOM	793	OG1	THR	98	-64.498	75.236	-19.855	1.00	68.81
5	ATOM	794	CG2	THR	98	-66.141	73.826	-18.696	1.00	56.15
	ATOM	795	C	THR	98	-62.655	74.965	-17.587	1.00	60.32
	ATOM	796	O	THR	98	-61.970	74.879	-16.560	1.00	65.03
	ATOM	797	N	LEU	99	-62.126	75.141	-18.798	1.00	54.94
	ATOM	798	CA	LEU	99	-60.677	75.237	-18.995	1.00	51.13
10	ATOM	799	CB	LEU	99	-60.249	74.285	-20.109	1.00	50.56
	ATOM	800	CG	LEU	99	-60.255	72.795	-19.775	1.00	52.80
	ATOM	801	CD1	LEU	99	-60.235	71.964	-21.056	1.00	48.38
	ATOM	802	CD2	LEU	99	-59.053	72.479	-18.898	1.00	50.80
	ATOM	803	C	LEU	99	-60.182	76.651	-19.313	1.00	50.20
15	ATOM	804	O	LEU	99	-59.874	76.970	-20.471	1.00	50.50
	ATOM	805	N	PRO	100	-60.063	77.508	-18.283	1.00	47.97
	ATOM	806	CD	PRO	100	-60.202	77.201	-16.851	1.00	45.68
	ATOM	807	CA	PRO	100	-59.608	78.889	-18.467	1.00	46.54
	ATOM	808	CB	PRO	100	-59.481	79.411	-17.035	1.00	45.50
20	ATOM	809	CG	PRO	100	-60.446	78.564	-16.270	1.00	45.80
	ATOM	810	C	PRO	100	-58.292	78.985	-19.220	1.00	46.78
	ATOM	811	O	PRO	100	-57.291	78.409	-18.812	1.00	52.13
	ATOM	812	N	ASN	101	-58.301	79.715	-20.322	1.00	42.71
	ATOM	813	CA	ASN	101	-57.108	79.917	-21.122	1.00	41.37
25	ATOM	814	CB	ASN	101	-56.781	78.660	-21.919	1.00	43.24
	ATOM	815	CG	ASN	101	-55.989	77.649	-21.116	1.00	45.93
	ATOM	816	OD1	ASN	101	-54.818	77.867	-20.801	1.00	45.33
	ATOM	817	ND2	ASN	101	-56.628	76.531	-20.773	1.00	48.11
	ATOM	818	C	ASN	101	-57.364	81.079	-22.062	1.00	42.66
30	ATOM	819	O	ASN	101	-57.716	80.882	-23.226	1.00	44.79
	ATOM	820	N	ASN	102	-57.198	82.300	-21.564	1.00	40.46
	ATOM	821	CA	ASN	102	-57.437	83.437	-22.422	1.00	37.90
	ATOM	822	CB	ASN	102	-58.532	84.320	-21.844	1.00	34.01
	ATOM	823	CG	ASN	102	-59.914	83.866	-22.272	1.00	32.44
35	ATOM	824	OD1	ASN	102	-60.118	83.480	-23.420	1.00	29.07
	ATOM	825	ND2	ASN	102	-60.869	83.915	-21.355	1.00	33.79
	ATOM	826	C	ASN	102	-56.264	84.289	-22.864	1.00	39.20
	ATOM	827	O	ASN	102	-55.965	84.337	-24.051	1.00	46.07
	ATOM	828	N	SER	103	-55.580	84.960	-21.958	1.00	35.42
40	ATOM	829	CA	SER	103	-54.468	85.816	-22.407	1.00	40.28
	ATOM	830	CB	SER	103	-53.462	85.038	-23.291	1.00	40.68
	ATOM	831	OG	SER	103	-53.619	85.310	-24.679	1.00	31.76
	ATOM	832	C	SER	103	-55.004	87.043	-23.182	1.00	35.24
	ATOM	833	O	SER	103	-55.714	86.925	-24.179	1.00	24.65
45	ATOM	834	N	CYS	104	-54.671	88.223	-22.676	1.00	32.88
	ATOM	835	CA	CYS	104	-55.097	89.461	-23.277	1.00	31.77
	ATOM	836	C	CYS	104	-53.847	90.322	-23.531	1.00	30.58
	ATOM	837	O	CYS	104	-52.963	90.420	-22.684	1.00	29.23
	ATOM	838	CB	CYS	104	-56.107	90.197	-22.353	1.00	28.43
50	ATOM	839	SG	CYS	104	-56.294	91.881	-22.961	1.00	46.07
	ATOM	840	N	TYR	105	-53.764	90.917	-24.720	1.00	30.34
	ATOM	841	CA	TYR	105	-52.656	91.796	-25.087	1.00	26.12
	ATOM	842	CB	TYR	105	-52.077	91.396	-26.443	1.00	27.03
	ATOM	843	CG	TYR	105	-51.047	92.361	-27.008	1.00	25.61
55	ATOM	844	CD1	TYR	105	-49.675	92.128	-26.862	1.00	24.90
	ATOM	845	CE1	TYR	105	-48.735	93.009	-27.376	1.00	26.08
	ATOM	846	CD2	TYR	105	-51.441	93.510	-27.686	1.00	25.41
	ATOM	847	CE2	TYR	105	-50.505	94.398	-28.200	1.00	28.91
	ATOM	848	CZ	TYR	105	-49.155	94.141	-28.043	1.00	27.58
60	ATOM	849	OH	TYR	105	-48.230	95.018	-28.566	1.00	29.89
	ATOM	850	C	TYR	105	-53.190	93.224	-25.179	1.00	29.20
	ATOM	851	O	TYR	105	-54.318	93.447	-25.607	1.00	33.10
	ATOM	852	N	SER	106	-52.385	94.193	-24.778	1.00	30.48
	ATOM	853	CA	SER	106	-52.813	95.581	-24.846	1.00	29.22
65	ATOM	854	CB	SER	106	-53.660	95.929	-23.619	1.00	28.30
	ATOM	855	OG	SER	106	-54.139	97.258	-23.674	1.00	28.62

	ATOM	856	C	SER	106	-51.572	96.454	-24.901	1.00	29.94
	ATOM	857	O	SER	106	-50.574	96.140	-24.265	1.00	29.72
	ATOM	858	N	ALA	107	-51.627	97.533	-25.677	1.00	27.64
	ATOM	859	CA	ALA	107	-50.495	98.441	-25.803	1.00	24.33
5	ATOM	860	CB	ALA	107	-49.523	97.925	-26.849	1.00	21.30
	ATOM	861	C	ALA	107	-50.937	99.851	-26.159	1.00	25.56
	ATOM	862	O	ALA	107	-52.070	100.080	-26.562	1.00	28.03
	ATOM	863	N	GLY	108	-50.028	100.800	-26.007	1.00	26.07
	ATOM	864	CA	GLY	108	-50.336	102.178	-26.329	1.00	25.49
10	ATOM	865	C	GLY	108	-49.109	103.043	-26.158	1.00	29.03
	ATOM	866	O	GLY	108	-48.072	102.561	-25.725	1.00	32.20
	ATOM	867	N	ILE	109	-49.221	104.321	-26.495	1.00	29.34
	ATOM	868	CA	ILE	109	-48.104	105.244	-26.370	1.00	27.71
	ATOM	869	CB	ILE	109	-47.931	106.074	-27.653	1.00	27.48
15	ATOM	870	CG2	ILE	109	-46.779	107.044	-27.500	1.00	26.46
	ATOM	871	CG1	ILE	109	-47.685	105.137	-28.835	1.00	26.16
	ATOM	872	CD1	ILE	109	-47.668	105.827	-30.169	1.00	29.07
	ATOM	873	C	ILE	109	-48.341	106.181	-25.202	1.00	28.66
	ATOM	874	O	ILE	109	-49.470	106.597	-24.949	1.00	30.59
20	ATOM	875	N	ALA	110	-47.278	106.510	-24.481	1.00	25.60
	ATOM	876	CA	ALA	110	-47.378	107.409	-23.337	1.00	26.36
	ATOM	877	CB	ALA	110	-47.624	106.616	-22.076	1.00	20.26
	ATOM	878	C	ALA	110	-46.096	108.209	-23.195	1.00	29.61
	ATOM	879	O	ALA	110	-45.041	107.779	-23.638	1.00	27.61
25	ATOM	880	N	LYS	111	-46.182	109.387	-22.595	1.00	33.77
	ATOM	881	CA	LYS	111	-44.988	110.175	-22.399	1.00	35.09
	ATOM	882	CB	LYS	111	-45.264	111.665	-22.565	1.00	39.38
	ATOM	883	CG	LYS	111	-44.038	112.490	-22.195	1.00	49.74
	ATOM	884	CD	LYS	111	-43.889	113.734	-23.028	1.00	53.95
30	ATOM	885	CE	LYS	111	-42.559	114.408	-22.721	1.00	54.28
	ATOM	886	NZ	LYS	111	-42.402	115.668	-23.524	1.00	60.85
	ATOM	887	C	LYS	111	-44.461	109.892	-20.997	1.00	35.28
	ATOM	888	O	LYS	111	-45.187	110.008	-20.018	1.00	36.58
	ATOM	889	N	LEU	112	-43.193	109.510	-20.919	1.00	32.32
35	ATOM	890	CA	LEU	112	-42.562	109.188	-19.654	1.00	29.67
	ATOM	891	CB	LEU	112	-42.100	107.733	-19.662	1.00	27.04
	ATOM	892	CG	LEU	112	-43.134	106.698	-20.103	1.00	24.38
	ATOM	893	CD1	LEU	112	-42.487	105.331	-20.191	1.00	27.56
	ATOM	894	CD2	LEU	112	-44.286	106.683	-19.137	1.00	19.52
40	ATOM	895	C	LEU	112	-41.367	110.098	-19.416	1.00	32.58
	ATOM	896	O	LEU	112	-40.890	110.762	-20.336	1.00	31.51
	ATOM	897	N	GLU	113	-40.882	110.122	-18.181	1.00	35.75
	ATOM	898	CA	GLU	113	-39.745	110.954	-17.829	1.00	40.08
	ATOM	899	CB	GLU	113	-40.177	112.130	-16.974	1.00	44.23
45	ATOM	900	CG	GLU	113	-41.377	112.887	-17.463	1.00	50.12
	ATOM	901	CD	GLU	113	-41.404	114.279	-16.875	1.00	58.08
	ATOM	902	OE1	GLU	113	-41.037	114.428	-15.672	1.00	61.13
	ATOM	903	OE2	GLU	113	-41.789	115.219	-17.617	1.00	63.64
	ATOM	904	C	GLU	113	-38.711	110.192	-17.038	1.00	41.63
50	ATOM	905	O	GLU	113	-39.006	109.160	-16.450	1.00	48.29
	ATOM	906	N	GLU	114	-37.494	110.726	-17.023	1.00	40.97
	ATOM	907	CA	GLU	114	-36.389	110.147	-16.271	1.00	36.68
	ATOM	908	CB	GLU	114	-35.260	111.147	-16.147	1.00	41.19
	ATOM	909	CG	GLU	114	-34.156	111.033	-17.135	1.00	46.58
55	ATOM	910	CD	GLU	114	-32.874	111.594	-16.557	1.00	51.12
	ATOM	911	OE1	GLU	114	-32.354	110.987	-15.589	1.00	51.88
	ATOM	912	OE2	GLU	114	-32.400	112.643	-17.057	1.00	57.08
	ATOM	913	C	GLU	114	-36.859	109.859	-14.861	1.00	35.51
	ATOM	914	O	GLU	114	-37.307	110.763	-14.166	1.00	38.92
60	ATOM	915	N	GLY	115	-36.744	108.614	-14.427	1.00	32.99
	ATOM	916	CA	GLY	115	-37.155	108.302	-13.076	1.00	31.79
	ATOM	917	C	GLY	115	-38.503	107.640	-12.981	1.00	34.39
	ATOM	918	O	GLY	115	-38.824	107.062	-11.952	1.00	39.20
	ATOM	919	N	ASP	116	-39.304	107.743	-14.032	1.00	35.66
65	ATOM	920	CA	ASP	116	-40.608	107.104	-14.028	1.00	38.09
	ATOM	921	CB	ASP	116	-41.424	107.492	-15.267	1.00	36.00

	ATOM	922	CG	ASP	116	-42.022	108.878	-15.173	1.00	35.09
	ATOM	923	OD1	ASP	116	-42.073	109.446	-14.062	1.00	33.81
	ATOM	924	OD2	ASP	116	-42.463	109.396	-16.216	1.00	35.24
	ATOM	925	C	ASP	116	-40.377	105.600	-14.045	1.00	39.38
5	ATOM	926	O	ASP	116	-39.347	105.126	-14.532	1.00	37.91
	ATOM	927	N	GLU	117	-41.326	104.849	-13.501	1.00	41.54
	ATOM	928	CA	GLU	117	-41.218	103.396	-13.504	1.00	37.92
	ATOM	929	CB	GLU	117	-41.027	102.860	-12.091	1.00	38.90
	ATOM	930	CG	GLU	117	-39.814	103.408	-11.385	1.00	48.15
10	ATOM	931	CD	GLU	117	-39.642	102.815	-10.002	1.00	52.62
	ATOM	932	OE1	GLU	117	-40.673	102.592	-9.310	1.00	55.49
	ATOM	933	OE2	GLU	117	-38.473	102.583	-9.608	1.00	57.10
	ATOM	934	C	GLU	117	-42.499	102.832	-14.087	1.00	34.02
	ATOM	935	O	GLU	117	-43.575	103.390	-13.880	1.00	33.31
15	ATOM	936	N	LEU	118	-42.377	101.746	-14.837	1.00	30.68
	ATOM	937	CA	LEU	118	-43.533	101.090	-15.424	1.00	27.14
	ATOM	938	CB	LEU	118	-43.323	100.871	-16.922	1.00	26.49
	ATOM	939	CG	LEU	118	-43.227	102.093	-17.828	1.00	26.21
	ATOM	940	CD1	LEU	118	-42.935	101.646	-19.242	1.00	23.05
20	ATOM	941	CD2	LEU	118	-44.520	102.880	-17.775	1.00	25.38
	ATOM	942	C	LEU	118	-43.708	99.736	-14.748	1.00	28.33
	ATOM	943	O	LEU	118	-42.725	99.088	-14.383	1.00	31.46
	ATOM	944	N	GLN	119	-44.953	99.308	-14.565	1.00	28.96
	ATOM	945	CA	GLN	119	-45.207	98.005	-13.965	1.00	29.13
25	ATOM	946	CB	GLN	119	-45.265	98.123	-12.449	1.00	28.23
	ATOM	947	CG	GLN	119	-46.472	98.848	-11.936	1.00	36.38
	ATOM	948	CD	GLN	119	-46.452	99.001	-10.435	1.00	36.57
	ATOM	949	OE1	GLN	119	-47.477	99.305	-9.824	1.00	40.70
	ATOM	950	NE2	GLN	119	-45.288	98.800	-9.829	1.00	36.80
30	ATOM	951	C	GLN	119	-46.489	97.371	-14.495	1.00	29.43
	ATOM	952	O	GLN	119	-47.403	98.066	-14.933	1.00	28.80
	ATOM	953	N	LEU	120	-46.527	96.043	-14.465	1.00	32.26
	ATOM	954	CA	LEU	120	-47.664	95.254	-14.929	1.00	29.96
	ATOM	955	CB	LEU	120	-47.165	94.134	-15.854	1.00	31.17
35	ATOM	956	CG	LEU	120	-48.058	93.265	-16.748	1.00	30.26
	ATOM	957	CD1	LEU	120	-49.255	92.749	-15.990	1.00	27.10
	ATOM	958	CD2	LEU	120	-48.486	94.076	-17.938	1.00	30.96
	ATOM	959	C	LEU	120	-48.318	94.651	-13.684	1.00	30.51
	ATOM	960	O	LEU	120	-47.725	93.814	-13.002	1.00	32.88
40	ATOM	961	N	ALA	121	-49.543	95.074	-13.394	1.00	27.40
	ATOM	962	CA	ALA	121	-50.255	94.588	-12.220	1.00	25.55
	ATOM	963	CB	ALA	121	-50.505	95.746	-11.267	1.00	19.41
	ATOM	964	C	ALA	121	-51.577	93.885	-12.531	1.00	29.89
	ATOM	965	O	ALA	121	-52.343	94.323	-13.387	1.00	35.88
45	ATOM	966	N	ILE	122	-51.828	92.784	-11.828	1.00	30.33
	ATOM	967	CA	ILE	122	-53.064	92.030	-11.989	1.00	30.27
	ATOM	968	CB	ILE	122	-52.796	90.529	-12.131	1.00	29.27
	ATOM	969	CG2	ILE	122	-54.100	89.808	-12.387	1.00	30.38
	ATOM	970	CG1	ILE	122	-51.819	90.280	-13.283	1.00	26.03
50	ATOM	971	CD1	ILE	122	-51.451	88.824	-13.499	1.00	29.21
	ATOM	972	C	ILE	122	-53.873	92.280	-10.720	1.00	33.20
	ATOM	973	O	ILE	122	-53.465	91.882	-9.632	1.00	33.30
	ATOM	974	N	PRO	123	-55.030	92.954	-10.846	1.00	34.70
	ATOM	975	CD	PRO	123	-55.565	93.497	-12.099	1.00	34.40
55	ATOM	976	CA	PRO	123	-55.924	93.287	-9.729	1.00	36.49
	ATOM	977	CB	PRO	123	-56.882	94.316	-10.336	1.00	30.22
	ATOM	978	CG	PRO	123	-56.217	94.748	-11.611	1.00	36.72
	ATOM	979	C	PRO	123	-56.692	92.095	-9.161	1.00	39.64
	ATOM	980	O	PRO	123	-57.926	92.116	-9.119	1.00	40.48
60	ATOM	981	N	ARG	124	-55.969	91.066	-8.728	1.00	44.58
	ATOM	982	CA	ARG	124	-56.593	89.876	-8.168	1.00	48.04
	ATOM	983	CB	ARG	124	-56.898	88.868	-9.270	1.00	52.91
	ATOM	984	CG	ARG	124	-57.735	89.433	-10.403	1.00	60.50
	ATOM	985	CD	ARG	124	-59.123	88.835	-10.332	1.00	72.57
65	ATOM	986	NE	ARG	124	-60.043	89.458	-11.300	1.00	84.90
	ATOM	987	CZ	ARG	124	-60.825	90.517	-11.054	1.00	85.44

	ATOM	988	NH1	ARG	124	-60.810	91.098	-9.853	1.00	87.69
	ATOM	989	NH2	ARG	124	-61.622	91.000	-12.017	1.00	82.78
	ATOM	990	C	ARG	124	-55.721	89.228	-7.105	1.00	48.39
	ATOM	991	O	ARG	124	-54.493	89.277	-7.187	1.00	45.26
5	ATOM	992	N	GLU	125	-56.378	88.619	-6.121	1.00	52.84
	ATOM	993	CA	GLU	125	-55.717	87.961	-4.996	1.00	55.02
	ATOM	994	CB	GLU	125	-56.758	87.217	-4.155	1.00	63.62
	ATOM	995	CG	GLU	125	-57.968	86.723	-4.953	1.00	70.62
	ATOM	996	CD	GLU	125	-58.658	87.842	-5.723	1.00	74.76
10	ATOM	997	OE1	GLU	125	-59.215	88.766	-5.071	1.00	75.75
	ATOM	998	OE2	GLU	125	-58.640	87.796	-6.979	1.00	80.71
	ATOM	999	C	GLU	125	-54.584	87.016	-5.370	1.00	53.44
	ATOM	1000	O	GLU	125	-53.419	87.313	-5.102	1.00	57.50
	ATOM	1001	N	ASN	126	-54.906	85.863	-5.940	1.00	47.18
15	ATOM	1002	CA	ASN	126	-53.854	84.942	-6.348	1.00	47.59
	ATOM	1003	CB	ASN	126	-53.787	83.717	-5.439	1.00	51.03
	ATOM	1004	CG	ASN	126	-52.654	83.810	-4.428	1.00	57.07
	ATOM	1005	OD1	ASN	126	-52.740	84.542	-3.430	1.00	61.15
	ATOM	1006	ND2	ASN	126	-51.571	83.079	-4.693	1.00	56.63
20	ATOM	1007	C	ASN	126	-54.136	84.525	-7.766	1.00	47.06
	ATOM	1008	O	ASN	126	-54.509	83.382	-8.044	1.00	45.61
	ATOM	1009	N	ALA	127	-53.960	85.488	-8.660	1.00	43.50
	ATOM	1010	CA	ALA	127	-54.212	85.291	-10.070	1.00	38.26
	ATOM	1011	CB	ALA	127	-53.658	86.461	-10.848	1.00	37.97
25	ATOM	1012	C	ALA	127	-53.618	83.996	-10.589	1.00	37.16
	ATOM	1013	O	ALA	127	-52.440	83.710	-10.383	1.00	33.14
	ATOM	1014	N	GLN	128	-54.456	83.201	-11.245	1.00	37.87
	ATOM	1015	CA	GLN	128	-53.997	81.958	-11.847	1.00	37.19
	ATOM	1016	CB	GLN	128	-55.149	80.962	-11.917	1.00	40.23
30	ATOM	1017	CG	GLN	128	-55.568	80.481	-10.548	1.00	38.14
	ATOM	1018	CD	GLN	128	-54.393	79.923	-9.782	1.00	39.51
	ATOM	1019	OE1	GLN	128	-53.843	78.874	-10.137	1.00	40.36
	ATOM	1020	NE2	GLN	128	-53.978	80.632	-8.733	1.00	41.53
	ATOM	1021	C	GLN	128	-53.509	82.362	-13.235	1.00	35.55
35	ATOM	1022	O	GLN	128	-54.290	82.554	-14.172	1.00	35.57
	ATOM	1023	N	ILE	129	-52.195	82.497	-13.337	1.00	34.64
	ATOM	1024	CA	ILE	129	-51.526	82.962	-14.544	1.00	29.60
	ATOM	1025	CB	ILE	129	-50.857	84.344	-14.208	1.00	31.03
	ATOM	1026	CG2	ILE	129	-49.430	84.423	-14.688	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-51.714	85.462	-14.756	1.00	26.37
	ATOM	1028	CD1	ILE	129	-53.075	85.482	-14.168	1.00	34.70
	ATOM	1029	C	ILE	129	-50.491	81.974	-15.072	1.00	29.55
	ATOM	1030	O	ILE	129	-49.955	81.171	-14.322	1.00	33.42
	ATOM	1031	N	SER	130	-50.215	82.025	-16.370	1.00	27.67
45	ATOM	1032	CA	SER	130	-49.195	81.159	-16.952	1.00	25.00
	ATOM	1033	CB	SER	130	-49.539	80.789	-18.385	1.00	22.09
	ATOM	1034	OG	SER	130	-48.429	80.150	-19.000	1.00	20.89
	ATOM	1035	C	SER	130	-47.882	81.926	-16.951	1.00	26.03
	ATOM	1036	O	SER	130	-47.860	83.094	-17.298	1.00	30.77
50	ATOM	1037	N	LEU	131	-46.791	81.279	-16.573	1.00	24.49
	ATOM	1038	CA	LEU	131	-45.504	81.957	-16.532	1.00	28.07
	ATOM	1039	CB	LEU	131	-44.835	81.740	-15.175	1.00	28.87
	ATOM	1040	CG	LEU	131	-45.178	82.745	-14.076	1.00	30.76
	ATOM	1041	CD1	LEU	131	-46.660	82.966	-13.990	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-44.644	82.231	-12.765	1.00	33.37
	ATOM	1043	C	LEU	131	-44.543	81.565	-17.641	1.00	33.66
	ATOM	1044	O	LEU	131	-43.327	81.587	-17.449	1.00	37.67
	ATOM	1045	N	ASP	132	-45.085	81.206	-18.799	1.00	34.97
	ATOM	1046	CA	ASP	132	-44.254	80.843	-19.943	1.00	34.31
60	ATOM	1047	CB	ASP	132	-45.021	79.943	-20.911	1.00	42.68
	ATOM	1048	CG	ASP	132	-45.138	78.512	-20.413	1.00	47.98
	ATOM	1049	OD1	ASP	132	-45.920	77.735	-21.014	1.00	50.41
	ATOM	1050	OD2	ASP	132	-44.438	78.167	-19.428	1.00	46.42
	ATOM	1051	C	ASP	132	-43.827	82.120	-20.647	1.00	34.32
65	ATOM	1052	O	ASP	132	-44.639	83.005	-20.895	1.00	32.87
	ATOM	1053	N	GLY	133	-42.541	82.204	-20.962	1.00	34.53

	ATOM	1054	CA	GLY	133	-42.007	83.386	-21.608	1.00	36.93
	ATOM	1055	C	GLY	133	-42.677	83.843	-22.888	1.00	35.06
	ATOM	1056	O	GLY	133	-42.568	85.010	-23.262	1.00	40.44
	ATOM	1057	N	ASP	134	-43.376	82.944	-23.566	1.00	31.08
5	ATOM	1058	CA	ASP	134	-44.015	83.320	-24.812	1.00	27.63
	ATOM	1059	CB	ASP	134	-43.920	82.170	-25.818	1.00	28.43
	ATOM	1060	CG	ASP	134	-44.593	80.903	-25.337	1.00	31.78
	ATOM	1061	OD1	ASP	134	-44.574	80.639	-24.120	1.00	35.68
	ATOM	1062	OD2	ASP	134	-45.130	80.157	-26.182	1.00	32.53
10	ATOM	1063	C	ASP	134	-45.447	83.799	-24.659	1.00	25.08
	ATOM	1064	O	ASP	134	-45.948	84.500	-25.510	1.00	19.71
	ATOM	1065	N	VAL	135	-46.103	83.462	-23.560	1.00	26.84
	ATOM	1066	CA	VAL	135	-47.479	83.906	-23.391	1.00	23.72
	ATOM	1067	CB	VAL	135	-48.370	82.774	-22.874	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-48.547	81.752	-23.964	1.00	22.76
	ATOM	1069	CG2	VAL	135	-47.760	82.144	-21.641	1.00	22.12
	ATOM	1070	C	VAL	135	-47.667	85.142	-22.518	1.00	26.54
	ATOM	1071	O	VAL	135	-48.655	85.848	-22.670	1.00	29.26
	ATOM	1072	N	THR	136	-46.743	85.412	-21.598	1.00	26.39
20	ATOM	1073	CA	THR	136	-46.871	86.619	-20.784	1.00	28.40
	ATOM	1074	CB	THR	136	-47.394	86.305	-19.342	1.00	28.21
	ATOM	1075	OG1	THR	136	-46.320	86.354	-18.406	1.00	37.13
	ATOM	1076	CG2	THR	136	-48.040	84.944	-19.292	1.00	26.90
	ATOM	1077	C	THR	136	-45.546	87.395	-20.754	1.00	31.60
25	ATOM	1078	O	THR	136	-44.520	86.901	-20.277	1.00	27.29
	ATOM	1079	N	PHE	137	-45.585	88.613	-21.298	1.00	31.45
	ATOM	1080	CA	PHE	137	-44.406	89.464	-21.395	1.00	26.64
	ATOM	1081	CB	PHE	137	-43.742	89.224	-22.743	1.00	26.84
	ATOM	1082	CG	PHE	137	-44.710	89.133	-23.886	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-45.145	90.279	-24.544	1.00	27.77
	ATOM	1084	CD2	PHE	137	-45.181	87.903	-24.315	1.00	23.70
	ATOM	1085	CE1	PHE	137	-46.028	90.201	-25.607	1.00	27.32
	ATOM	1086	CE2	PHE	137	-46.067	87.817	-25.378	1.00	26.31
	ATOM	1087	CZ	PHE	137	-46.489	88.970	-26.024	1.00	29.43
35	ATOM	1088	C	PHE	137	-44.731	90.942	-21.194	1.00	28.06
	ATOM	1089	O	PHE	137	-45.889	91.331	-21.246	1.00	25.04
	ATOM	1090	N	PHE	138	-43.702	91.759	-20.973	1.00	29.81
	ATOM	1091	CA	PHE	138	-43.889	93.182	-20.700	1.00	29.93
	ATOM	1092	CB	PHE	138	-43.615	93.408	-19.214	1.00	27.47
40	ATOM	1093	CG	PHE	138	-44.038	94.744	-18.706	1.00	28.49
	ATOM	1094	CD1	PHE	138	-45.026	95.479	-19.352	1.00	27.65
	ATOM	1095	CD2	PHE	138	-43.444	95.272	-17.562	1.00	31.06
	ATOM	1096	CE1	PHE	138	-45.412	96.721	-18.867	1.00	30.90
	ATOM	1097	CE2	PHE	138	-43.821	96.514	-17.066	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-44.807	97.241	-17.719	1.00	27.41
	ATOM	1099	C	PHE	138	-43.042	94.110	-21.593	1.00	30.91
	ATOM	1100	O	PHE	138	-41.830	93.965	-21.701	1.00	24.94
	ATOM	1101	N	GLY	139	-43.736	95.081	-22.193	1.00	36.94
	ATOM	1102	CA	GLY	139	-43.208	96.039	-23.163	1.00	36.08
50	ATOM	1103	C	GLY	139	-42.127	97.082	-23.020	1.00	36.48
	ATOM	1104	O	GLY	139	-41.045	96.781	-22.551	1.00	43.61
	ATOM	1105	N	ALA	140	-42.419	98.289	-23.506	1.00	36.60
	ATOM	1106	CA	ALA	140	-41.516	99.456	-23.486	1.00	34.94
	ATOM	1107	CB	ALA	140	-40.810	99.552	-22.145	1.00	36.94
55	ATOM	1108	C	ALA	140	-40.480	99.596	-24.618	1.00	32.53
	ATOM	1109	O	ALA	140	-39.460	98.916	-24.643	1.00	27.96
	ATOM	1110	N	LEU	141	-40.756	100.517	-25.538	1.00	33.23
	ATOM	1111	CA	LEU	141	-39.888	100.815	-26.685	1.00	36.11
	ATOM	1112	CB	LEU	141	-40.431	100.127	-27.937	1.00	37.53
60	ATOM	1113	CG	LEU	141	-39.774	100.337	-29.305	1.00	38.87
	ATOM	1114	CD1	LEU	141	-40.356	99.342	-30.279	1.00	40.67
	ATOM	1115	CD2	LEU	141	-40.006	101.743	-29.822	1.00	37.53
	ATOM	1116	C	LEU	141	-39.872	102.333	-26.895	1.00	38.89
	ATOM	1117	O	LEU	141	-40.926	102.961	-26.970	1.00	39.25
65	ATOM	1118	N	LYS	142	-38.691	102.933	-27.001	1.00	42.86
	ATOM	1119	CA	LYS	142	-38.637	104.380	-27.182	1.00	42.96

	ATOM	1120	CB	LYS	142	-37.338	104.938	-26.609	1.00	41.80
	ATOM	1121	CG	LYS	142	-37.226	106.432	-26.800	1.00	46.88
	ATOM	1122	CD	LYS	142	-36.225	107.069	-25.858	1.00	49.58
	ATOM	1123	CE	LYS	142	-36.173	108.567	-26.110	1.00	51.07
5	ATOM	1124	NZ	LYS	142	-35.272	109.270	-25.165	1.00	56.20
	ATOM	1125	C	LYS	142	-38.805	104.857	-28.621	1.00	43.41
	ATOM	1126	O	LYS	142	-38.128	104.381	-29.527	1.00	44.98
	ATOM	1127	N	LEU	143	-39.712	105.809	-28.820	1.00	42.81
	ATOM	1128	CA	LEU	143	-39.972	106.362	-30.147	1.00	40.65
10	ATOM	1129	CB	LEU	143	-41.389	106.929	-30.218	1.00	36.56
	ATOM	1130	CG	LEU	143	-42.560	105.999	-29.903	1.00	36.17
	ATOM	1131	CD1	LEU	143	-43.854	106.788	-29.975	1.00	37.95
	ATOM	1132	CD2	LEU	143	-42.585	104.844	-30.881	1.00	27.38
	ATOM	1133	C	LEU	143	-38.987	107.484	-30.459	1.00	43.14
15	ATOM	1134	O	LEU	143	-38.537	108.195	-29.555	1.00	46.28
	ATOM	1135	N	LEU	144	-38.652	107.647	-31.732	1.00	42.63
	ATOM	1136	CA	LEU	144	-37.733	108.704	-32.124	1.00	44.11
	ATOM	1137	CB	LEU	144	-37.125	108.409	-33.492	1.00	44.88
	ATOM	1138	CG	LEU	144	-36.232	107.170	-33.532	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-35.770	106.915	-34.941	1.00	47.71
	ATOM	1140	CD2	LEU	144	-35.040	107.368	-32.613	1.00	47.87
	ATOM	1141	C	LEU	144	-38.467	110.029	-32.180	1.00	45.66
	ATOM	1142	O	LEU	144	-39.707	110.014	-32.292	1.00	46.37
	ATOM	1143	OXT	LEU	144	-37.787	111.071	-32.124	1.00	50.66
25	END					-94.358	153.908	54.620	0.00	0.00

TABLE 11

	111								
	ATOM	1	CB	VAL	1	-35.659	114.317	-37.002	1.00 69.28
5	ATOM	2	CG1	VAL	1	-34.610	114.343	-35.870	1.00 68.26
	ATOM	3	CG2	VAL	1	-35.166	115.085	-38.234	1.00 71.75
	ATOM	4	C	VAL	1	-36.851	112.236	-36.254	1.00 64.86
	ATOM	5	O	VAL	1	-36.354	111.432	-35.452	1.00 64.02
	ATOM	6	N	VAL	1	-36.727	112.773	-38.692	1.00 65.10
10	ATOM	7	CA	VAL	1	-36.000	112.841	-37.388	1.00 65.98
	ATOM	8	N	THR	2	-38.124	112.632	-36.185	1.00 61.42
	ATOM	9	CA	THR	2	-39.025	112.120	-35.152	1.00 56.64
	ATOM	10	CB	THR	2	-39.578	113.246	-34.261	1.00 56.77
	ATOM	11	OG1	THR	2	-40.209	114.229	-35.086	1.00 57.45
15	ATOM	12	CG2	THR	2	-38.466	113.899	-33.449	1.00 55.68
	ATOM	13	C	THR	2	-40.218	111.406	-35.755	1.00 53.08
	ATOM	14	O	THR	2	-40.460	111.475	-36.956	1.00 52.00
	ATOM	15	N	GLN	3	-40.978	110.730	-34.902	1.00 50.34
	ATOM	16	CA	GLN	3	-42.158	109.998	-35.344	1.00 45.97
20	ATOM	17	CB	GLN	3	-42.093	108.557	-34.853	1.00 46.09
	ATOM	18	CG	GLN	3	-40.776	107.878	-35.124	1.00 49.60
	ATOM	19	CD	GLN	3	-40.788	106.433	-34.695	1.00 50.43
	ATOM	20	OE1	GLN	3	-41.491	105.616	-35.280	1.00 50.94
	ATOM	21	NE2	GLN	3	-40.014	106.109	-33.661	1.00 49.99
25	ATOM	22	C	GLN	3	-43.426	110.647	-34.820	1.00 42.41
	ATOM	23	O	GLN	3	-43.716	110.602	-33.628	1.00 37.34
	ATOM	24	N	ASP	4	-44.182	111.254	-35.720	1.00 40.32
	ATOM	25	CA	ASP	4	-45.420	111.906	-35.334	1.00 39.37
	ATOM	26	CB	ASP	4	-45.981	112.710	-36.510	1.00 44.56
30	ATOM	27	CG	ASP	4	-45.089	113.870	-36.899	1.00 45.76
	ATOM	28	OD1	ASP	4	-43.894	113.845	-36.525	1.00 48.58
	ATOM	29	OD2	ASP	4	-45.577	114.796	-37.581	1.00 45.55
	ATOM	30	C	ASP	4	-46.423	110.854	-34.915	1.00 36.30
	ATOM	31	O	ASP	4	-46.431	109.744	-35.442	1.00 36.29
35	ATOM	32	N	CYS	5	-47.263	111.203	-33.954	1.00 35.11
	ATOM	33	CA	CYS	5	-48.297	110.297	-33.487	1.00 35.61
	ATOM	34	CB	CYS	5	-47.712	109.214	-32.576	1.00 34.96
	ATOM	35	SG	CYS	5	-46.506	109.782	-31.395	1.00 35.34
	ATOM	36	C	CYS	5	-49.375	111.081	-32.771	1.00 31.80
40	ATOM	37	O	CYS	5	-49.119	112.143	-32.215	1.00 31.85
	ATOM	38	N	LEU	6	-50.593	110.566	-32.824	1.00 27.72
	ATOM	39	CA	LEU	6	-51.725	111.198	-32.177	1.00 30.80
	ATOM	40	CB	LEU	6	-52.559	111.961	-33.199	1.00 29.25
	ATOM	41	CG	LEU	6	-53.843	112.633	-32.710	1.00 29.47
45	ATOM	42	CD1	LEU	6	-54.187	113.787	-33.614	1.00 30.65
	ATOM	43	CD2	LEU	6	-54.965	111.640	-32.703	1.00 35.27
	ATOM	44	C	LEU	6	-52.547	110.095	-31.539	1.00 31.83
	ATOM	45	O	LEU	6	-52.897	109.125	-32.199	1.00 33.45
	ATOM	46	N	GLN	7	-52.845	110.235	-30.253	1.00 32.23
50	ATOM	47	CA	GLN	7	-53.612	109.227	-29.553	1.00 30.26
	ATOM	48	CB	GLN	7	-52.780	108.649	-28.415	1.00 29.56
	ATOM	49	CG	GLN	7	-53.341	107.390	-27.790	1.00 25.09
	ATOM	50	CD	GLN	7	-52.344	106.736	-26.855	1.00 26.50
	ATOM	51	OE1	GLN	7	-52.055	107.246	-25.772	1.00 24.40
55	ATOM	52	NE2	GLN	7	-51.798	105.611	-27.278	1.00 27.03
	ATOM	53	C	GLN	7	-54.897	109.821	-29.020	1.00 32.11
	ATOM	54	O	GLN	7	-54.925	110.952	-28.552	1.00 34.24
	ATOM	55	N	LEU	8	-55.966	109.044	-29.117	1.00 33.73
	ATOM	56	CA	LEU	8	-57.280	109.458	-28.652	1.00 35.41
60	ATOM	57	CB	LEU	8	-58.291	109.394	-29.810	1.00 32.44
	ATOM	58	CG	LEU	8	-58.577	110.632	-30.672	1.00 27.69
	ATOM	59	CD1	LEU	8	-57.485	111.663	-30.549	1.00 26.43
	ATOM	60	CD2	LEU	8	-58.758	110.187	-32.098	1.00 18.64
	ATOM	61	C	LEU	8	-57.734	108.568	-27.505	1.00 36.87
65	ATOM	62	O	LEU	8	-57.415	107.381	-27.451	1.00 36.05
	ATOM	63	N	ILE	9	-58.493	109.158	-26.594	1.00 39.25

	ATOM	64	CA	ILE	9	-58.996	108.465	-25.420	1.00	35.57
	ATOM	65	CB	ILE	9	-58.309	109.055	-24.163	1.00	34.28
	ATOM	66	CG2	ILE	9	-59.301	109.330	-23.061	1.00	38.20
	ATOM	67	CG1	ILE	9	-57.226	108.114	-23.690	1.00	35.22
5	ATOM	68	CD1	ILE	9	-56.574	108.611	-22.445	1.00	43.63
	ATOM	69	C	ILE	9	-60.518	108.619	-25.348	1.00	35.28
	ATOM	70	O	ILE	9	-61.062	109.639	-25.761	1.00	33.75
	ATOM	71	N	ALA	10	-61.209	107.605	-24.842	1.00	35.78
	ATOM	72	CA	ALA	10	-62.658	107.688	-24.733	1.00	35.28
10	ATOM	73	CB	ALA	10	-63.223	106.377	-24.237	1.00	28.47
	ATOM	74	C	ALA	10	-63.057	108.815	-23.789	1.00	36.49
	ATOM	75	O	ALA	10	-62.441	109.018	-22.747	1.00	37.41
	ATOM	76	N	ASP	11	-64.095	109.549	-24.163	1.00	39.44
	ATOM	77	CA	ASP	11	-64.587	110.653	-23.348	1.00	40.77
15	ATOM	78	CB	ASP	11	-64.828	111.883	-24.224	1.00	38.90
	ATOM	79	CG	ASP	11	-65.462	113.019	-23.459	1.00	40.34
	ATOM	80	OD1	ASP	11	-65.284	113.084	-22.218	1.00	39.50
	ATOM	81	OD2	ASP	11	-66.129	113.851	-24.103	1.00	37.38
	ATOM	82	C	ASP	11	-65.875	110.247	-22.640	1.00	42.25
20	ATOM	83	O	ASP	11	-66.971	110.367	-23.189	1.00	40.54
	ATOM	84	N	SER	12	-65.725	109.772	-21.410	1.00	42.92
	ATOM	85	CA	SER	12	-66.849	109.299	-20.605	1.00	43.84
	ATOM	86	CB	SER	12	-66.328	108.689	-19.309	1.00	42.13
	ATOM	87	OG	SER	12	-65.566	109.640	-18.586	1.00	43.31
25	ATOM	88	C	SER	12	-67.891	110.352	-20.267	1.00	46.10
	ATOM	89	O	SER	12	-68.922	110.041	-19.671	1.00	43.90
	ATOM	90	N	GLU	13	-67.629	111.596	-20.652	1.00	46.72
	ATOM	91	CA	GLU	13	-68.560	112.668	-20.358	1.00	44.93
	ATOM	92	CB	GLU	13	-67.806	113.869	-19.816	1.00	46.71
30	ATOM	93	CG	GLU	13	-67.199	113.584	-18.470	1.00	58.25
	ATOM	94	CD	GLU	13	-66.979	114.846	-17.662	1.00	66.70
	ATOM	95	OE1	GLU	13	-67.986	115.568	-17.422	1.00	74.10
	ATOM	96	OE2	GLU	13	-65.813	115.118	-17.267	1.00	71.55
	ATOM	97	C	GLU	13	-69.485	113.087	-21.496	1.00	43.24
35	ATOM	98	O	GLU	13	-70.131	114.135	-21.422	1.00	44.25
	ATOM	99	N	THR	14	-69.546	112.286	-22.549	1.00	37.06
	ATOM	100	CA	THR	14	-70.464	112.580	-23.642	1.00	36.77
	ATOM	101	CB	THR	14	-69.785	113.331	-24.826	1.00	33.39
	ATOM	102	OG1	THR	14	-68.967	112.430	-25.572	1.00	41.69
40	ATOM	103	CG2	THR	14	-68.932	114.479	-24.312	1.00	31.66
	ATOM	104	C	THR	14	-71.018	111.238	-24.109	1.00	34.96
	ATOM	105	O	THR	14	-70.355	110.213	-24.002	1.00	35.04
	ATOM	106	N	PRO	15	-72.252	111.228	-24.614	1.00	37.11
	ATOM	107	CD	PRO	15	-73.139	112.391	-24.758	1.00	36.50
45	ATOM	108	CA	PRO	15	-72.909	110.006	-25.091	1.00	38.81
	ATOM	109	CB	PRO	15	-74.316	110.486	-25.445	1.00	39.15
	ATOM	110	CG	PRO	15	-74.483	111.751	-24.639	1.00	37.32
	ATOM	111	C	PRO	15	-72.216	109.392	-26.301	1.00	37.78
	ATOM	112	O	PRO	15	-71.590	110.103	-27.081	1.00	35.91
50	ATOM	113	N	THR	16	-72.327	108.078	-26.465	1.00	36.67
	ATOM	114	CA	THR	16	-71.710	107.443	-27.618	1.00	38.46
	ATOM	115	CB	THR	16	-71.788	105.906	-27.546	1.00	38.61
	ATOM	116	OG1	THR	16	-73.158	105.492	-27.577	1.00	42.44
	ATOM	117	CG2	THR	16	-71.138	105.403	-26.267	1.00	40.11
55	ATOM	118	C	THR	16	-72.504	107.911	-28.819	1.00	35.97
	ATOM	119	O	THR	16	-73.724	107.981	-28.754	1.00	39.85
	ATOM	120	N	ILE	17	-71.819	108.237	-29.907	1.00	33.66
	ATOM	121	CA	ILE	17	-72.492	108.711	-31.107	1.00	34.76
	ATOM	122	CB	ILE	17	-71.504	109.424	-32.037	1.00	32.44
60	ATOM	123	CG2	ILE	17	-72.189	109.804	-33.335	1.00	28.66
	ATOM	124	CG1	ILE	17	-70.932	110.656	-31.326	1.00	33.15
	ATOM	125	CD1	ILE	17	-69.878	111.405	-32.117	1.00	31.19
	ATOM	126	C	ILE	17	-73.187	107.604	-31.888	1.00	40.64
	ATOM	127	O	ILE	17	-72.583	106.591	-32.217	1.00	44.07
65	ATOM	128	N	GLN	18	-74.469	107.801	-32.185	1.00	45.91
	ATOM	129	CA	GLN	18	-75.241	106.820	-32.943	1.00	50.22

	ATOM	130	CB	GLN	18	-76.604	106.588	-32.298	1.00	53.09
	ATOM	131	CG	GLN	18	-76.842	105.143	-31.946	1.00	57.60
	ATOM	132	CD	GLN	18	-75.936	104.702	-30.827	1.00	61.39
	ATOM	133	OE1	GLN	18	-76.088	105.138	-29.683	1.00	68.36
5	ATOM	134	NE2	GLN	18	-74.971	103.851	-31.148	1.00	64.93
	ATOM	135	C	GLN	18	-75.456	107.294	-34.369	1.00	51.56
	ATOM	136	O	GLN	18	-75.844	108.435	-34.602	1.00	56.44
	ATOM	137	N	LYS	19	-75.212	106.416	-35.327	1.00	52.60
	ATOM	138	CA	LYS	19	-75.395	106.788	-36.720	1.00	56.10
10	ATOM	139	CB	LYS	19	-74.334	107.804	-37.148	1.00	55.55
	ATOM	140	CG	LYS	19	-74.429	108.129	-38.621	1.00	61.15
	ATOM	141	CD	LYS	19	-73.545	109.288	-39.033	1.00	64.42
	ATOM	142	CE	LYS	19	-73.710	109.569	-40.534	1.00	64.84
	ATOM	143	NZ	LYS	19	-72.887	110.731	-41.001	1.00	69.52
15	ATOM	144	C	LYS	19	-75.353	105.591	-37.657	1.00	57.41
	ATOM	145	O	LYS	19	-74.477	104.732	-37.538	1.00	61.02
	ATOM	146	N	GLY	20	-76.302	105.549	-38.592	1.00	58.84
	ATOM	147	CA	GLY	20	-76.372	104.462	-39.551	1.00	56.66
	ATOM	148	C	GLY	20	-76.281	103.094	-38.904	1.00	56.87
20	ATOM	149	O	GLY	20	-75.658	102.190	-39.464	1.00	58.64
	ATOM	150	N	SER	21	-76.901	102.943	-37.732	1.00	54.49
	ATOM	151	CA	SER	21	-76.896	101.681	-36.988	1.00	53.68
	ATOM	152	CB	SER	21	-77.562	100.571	-37.813	1.00	55.80
	ATOM	153	OG	SER	21	-76.651	99.961	-38.708	1.00	65.34
25	ATOM	154	C	SER	21	-75.474	101.250	-36.556	1.00	51.41
	ATOM	155	O	SER	21	-75.183	100.055	-36.372	1.00	47.74
	ATOM	156	N	TYR	22	-74.598	102.242	-36.403	1.00	46.25
	ATOM	157	CA	TYR	22	-73.224	102.032	-35.968	1.00	40.57
	ATOM	158	CB	TYR	22	-72.240	102.502	-37.041	1.00	43.27
30	ATOM	159	CG	TYR	22	-71.902	101.455	-38.071	1.00	49.18
	ATOM	160	CD1	TYR	22	-72.891	100.880	-38.861	1.00	52.61
	ATOM	161	CE1	TYR	22	-72.583	99.899	-39.808	1.00	51.92
	ATOM	162	CD2	TYR	22	-70.591	101.026	-38.250	1.00	50.01
	ATOM	163	CE2	TYR	22	-70.272	100.050	-39.195	1.00	50.94
35	ATOM	164	CZ	TYR	22	-71.273	99.491	-39.968	1.00	50.78
	ATOM	165	OH	TYR	22	-70.959	98.524	-40.898	1.00	53.87
	ATOM	166	C	TYR	22	-73.011	102.853	-34.700	1.00	37.85
	ATOM	167	O	TYR	22	-73.630	103.902	-34.527	1.00	38.70
	ATOM	168	N	THR	23	-72.156	102.375	-33.806	1.00	33.81
40	ATOM	169	CA	THR	23	-71.872	103.117	-32.583	1.00	32.40
	ATOM	170	CB	THR	23	-71.932	102.228	-31.315	1.00	31.58
	ATOM	171	OG1	THR	23	-73.213	101.598	-31.216	1.00	30.76
	ATOM	172	CG2	THR	23	-71.698	103.070	-30.079	1.00	24.08
	ATOM	173	C	THR	23	-70.463	103.672	-32.697	1.00	33.28
45	ATOM	174	O	THR	23	-69.538	102.952	-33.054	1.00	30.44
	ATOM	175	N	PHE	24	-70.306	104.953	-32.397	1.00	32.16
	ATOM	176	CA	PHE	24	-69.000	105.586	-32.460	1.00	30.68
	ATOM	177	CB	PHE	24	-69.006	106.728	-33.476	1.00	28.20
	ATOM	178	CG	PHE	24	-69.196	106.275	-34.888	1.00	31.00
50	ATOM	179	CD1	PHE	24	-70.466	106.049	-35.394	1.00	31.69
	ATOM	180	CD2	PHE	24	-68.098	106.040	-35.710	1.00	31.42
	ATOM	181	CE1	PHE	24	-70.639	105.595	-36.696	1.00	32.57
	ATOM	182	CE2	PHE	24	-68.263	105.589	-37.009	1.00	25.01
	ATOM	183	CZ	PHE	24	-69.536	105.366	-37.501	1.00	30.12
55	ATOM	184	C	PHE	24	-68.563	106.113	-31.099	1.00	31.73
	ATOM	185	O	PHE	24	-69.307	106.812	-30.419	1.00	33.31
	ATOM	186	N	VAL	25	-67.349	105.762	-30.705	1.00	32.92
	ATOM	187	CA	VAL	25	-66.809	106.207	-29.439	1.00	30.84
	ATOM	188	CB	VAL	25	-65.488	105.488	-29.121	1.00	30.29
60	ATOM	189	CG1	VAL	25	-64.921	105.997	-27.819	1.00	28.13
	ATOM	190	CG2	VAL	25	-65.713	104.002	-29.058	1.00	29.53
	ATOM	191	C	VAL	25	-66.531	107.705	-29.487	1.00	36.08
	ATOM	192	O	VAL	25	-66.009	108.217	-30.476	1.00	35.82
	ATOM	193	N	PRO	26	-66.902	108.435	-28.426	1.00	38.03
65	ATOM	194	CD	PRO	26	-67.795	107.990	-27.345	1.00	38.28
	ATOM	195	CA	PRO	26	-66.680	109.886	-28.344	1.00	38.08

	ATOM	196	CB	PRO	26	-67.628	110.326	-27.226	1.00	37.62
	ATOM	197	CG	PRO	26	-68.622	109.214	-27.121	1.00	36.31
	ATOM	198	C	PRO	26	-65.221	110.065	-27.928	1.00	40.85
	ATOM	199	O	PRO	26	-64.853	109.733	-26.803	1.00	42.27
5	ATOM	200	N	TRP	27	-64.384	110.585	-28.814	1.00	40.86
	ATOM	201	CA	TRP	27	-62.976	110.727	-28.468	1.00	37.05
	ATOM	202	CB	TRP	27	-62.099	110.536	-29.708	1.00	33.29
	ATOM	203	CG	TRP	27	-62.250	109.198	-30.339	1.00	31.06
	ATOM	204	CD2	TRP	27	-61.969	107.932	-29.742	1.00	27.79
10	ATOM	205	CE2	TRP	27	-62.289	106.947	-30.690	1.00	29.28
	ATOM	206	CE3	TRP	27	-61.479	107.534	-28.497	1.00	27.06
	ATOM	207	CD1	TRP	27	-62.711	108.939	-31.587	1.00	31.07
	ATOM	208	NE1	TRP	27	-62.739	107.591	-31.809	1.00	32.70
	ATOM	209	CZ2	TRP	27	-62.137	105.581	-30.435	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-61.327	106.176	-28.243	1.00	26.93
	ATOM	211	CH2	TRP	27	-61.656	105.217	-29.209	1.00	28.08
	ATOM	212	C	TRP	27	-62.592	112.024	-27.793	1.00	38.50
	ATOM	213	O	TRP	27	-63.282	113.032	-27.902	1.00	39.07
	ATOM	214	N	LEU	28	-61.469	111.964	-27.087	1.00	40.19
20	ATOM	215	CA	LEU	28	-60.899	113.101	-26.382	1.00	39.45
	ATOM	216	CB	LEU	28	-61.141	112.971	-24.884	1.00	44.18
	ATOM	217	CG	LEU	28	-60.950	114.246	-24.066	1.00	46.83
	ATOM	218	CD1	LEU	28	-62.044	115.247	-24.450	1.00	45.15
	ATOM	219	CD2	LEU	28	-61.011	113.919	-22.571	1.00	49.15
25	ATOM	220	C	LEU	28	-59.406	112.993	-26.663	1.00	39.80
	ATOM	221	O	LEU	28	-58.859	111.900	-26.647	1.00	42.40
	ATOM	222	N	LEU	29	-58.744	114.106	-26.931	1.00	33.58
	ATOM	223	CA	LEU	29	-57.323	114.039	-27.213	1.00	29.21
	ATOM	224	CB	LEU	29	-56.788	115.402	-27.649	1.00	27.16
30	ATOM	225	CG	LEU	29	-55.277	115.429	-27.900	1.00	22.36
	ATOM	226	CD1	LEU	29	-54.964	114.713	-29.203	1.00	21.47
	ATOM	227	CD2	LEU	29	-54.779	116.849	-27.943	1.00	14.53
	ATOM	228	C	LEU	29	-56.512	113.557	-26.015	1.00	29.98
	ATOM	229	O	LEU	29	-56.614	114.094	-24.917	1.00	30.36
35	ATOM	230	N	SER	30	-55.707	112.529	-26.245	1.00	29.88
	ATOM	231	CA	SER	30	-54.842	111.993	-25.212	1.00	28.49
	ATOM	232	CB	SER	30	-54.576	110.505	-25.442	1.00	26.13
	ATOM	233	OG	SER	30	-53.660	110.009	-24.491	1.00	20.25
	ATOM	234	C	SER	30	-53.553	112.792	-25.350	1.00	31.61
40	ATOM	235	O	SER	30	-53.062	113.374	-24.387	1.00	35.14
	ATOM	236	N	PHE	31	-53.013	112.822	-26.564	1.00	32.12
	ATOM	237	CA	PHE	31	-51.803	113.576	-26.843	1.00	30.76
	ATOM	238	CB	PHE	31	-50.595	112.954	-26.132	1.00	27.55
	ATOM	239	CG	PHE	31	-49.897	111.881	-26.920	1.00	31.23
45	ATOM	240	CD1	PHE	31	-48.963	112.212	-27.895	1.00	32.00
	ATOM	241	CD2	PHE	31	-50.183	110.536	-26.697	1.00	27.89
	ATOM	242	CE1	PHE	31	-48.328	111.225	-28.633	1.00	29.70
	ATOM	243	CE2	PHE	31	-49.550	109.546	-27.432	1.00	26.15
	ATOM	244	CZ	PHE	31	-48.623	109.890	-28.401	1.00	28.30
50	ATOM	245	C	PHE	31	-51.562	113.631	-28.342	1.00	32.06
	ATOM	246	O	PHE	31	-51.957	112.734	-29.086	1.00	32.26
	ATOM	247	N	LYS	32	-50.925	114.703	-28.779	1.00	34.83
	ATOM	248	CA	LYS	32	-50.610	114.878	-30.182	1.00	37.13
	ATOM	249	CB	LYS	32	-51.466	115.994	-30.777	1.00	36.55
55	ATOM	250	CG	LYS	32	-51.152	116.283	-32.219	1.00	40.62
	ATOM	251	CD	LYS	32	-51.800	117.559	-32.695	1.00	44.56
	ATOM	252	CE	LYS	32	-51.203	117.953	-34.048	1.00	46.29
	ATOM	253	NZ	LYS	32	-51.870	119.140	-34.660	1.00	50.58
	ATOM	254	C	LYS	32	-49.129	115.236	-30.261	1.00	36.39
60	ATOM	255	O	LYS	32	-48.648	116.080	-29.504	1.00	38.69
	ATOM	256	N	ARG	33	-48.404	114.582	-31.162	1.00	37.90
	ATOM	257	CA	ARG	33	-46.979	114.843	-31.313	1.00	35.74
	ATOM	258	CB	ARG	33	-46.192	113.689	-30.702	1.00	36.79
	ATOM	259	CG	ARG	33	-44.690	113.775	-30.850	1.00	31.09
65	ATOM	260	CD	ARG	33	-44.028	112.722	-29.969	1.00	31.62
	ATOM	261	NE	ARG	33	-42.572	112.829	-29.932	1.00	38.16

	ATOM	262	CZ	ARG	33	-41.749	112.058	-30.632	1.00	39.15
	ATOM	263	NH1	ARG	33	-42.235	111.115	-31.426	1.00	43.41
	ATOM	264	NH2	ARG	33	-40.440	112.227	-30.542	1.00	40.65
	ATOM	265	C	ARG	33	-46.630	115.007	-32.787	1.00	37.71
5	ATOM	266	O	ARG	33	-46.829	114.090	-33.589	1.00	41.16
	ATOM	267	N	GLY	34	-46.117	116.175	-33.155	1.00	36.59
	ATOM	268	CA	GLY	34	-45.762	116.397	-34.541	1.00	36.72
	ATOM	269	C	GLY	34	-46.884	117.031	-35.337	1.00	40.01
	ATOM	270	O	GLY	34	-47.858	117.532	-34.777	1.00	39.85
10	ATOM	271	N	SER	35	-46.753	116.981	-36.655	1.00	38.74
	ATOM	272	CA	SER	35	-47.733	117.585	-37.546	1.00	40.23
	ATOM	273	CB	SER	35	-47.039	118.655	-38.378	1.00	43.02
	ATOM	274	OG	SER	35	-45.897	118.106	-39.035	1.00	48.05
	ATOM	275	C	SER	35	-48.459	116.626	-38.493	1.00	40.54
15	ATOM	276	O	SER	35	-49.502	116.982	-39.041	1.00	43.74
	ATOM	277	N	ALA	36	-47.918	115.426	-38.686	1.00	35.56
	ATOM	278	CA	ALA	36	-48.502	114.452	-39.601	1.00	29.62
	ATOM	279	CB	ALA	36	-47.606	113.233	-39.676	1.00	25.47
	ATOM	280	C	ALA	36	-49.937	114.019	-39.310	1.00	31.23
20	ATOM	281	O	ALA	36	-50.646	113.602	-40.217	1.00	29.56
	ATOM	282	N	LEU	37	-50.370	114.117	-38.058	1.00	34.09
	ATOM	283	CA	LEU	37	-51.717	113.694	-37.690	1.00	32.32
	ATOM	284	CB	LEU	37	-51.651	112.392	-36.896	1.00	26.47
	ATOM	285	CG	LEU	37	-51.008	111.221	-37.635	1.00	25.72
25	ATOM	286	CD1	LEU	37	-50.568	110.160	-36.657	1.00	24.61
	ATOM	287	CD2	LEU	37	-51.981	110.662	-38.646	1.00	27.53
	ATOM	288	C	LEU	37	-52.470	114.739	-36.885	1.00	35.16
	ATOM	289	O	LEU	37	-51.900	115.425	-36.044	1.00	36.47
	ATOM	290	N	GLU	38	-53.763	114.845	-37.149	1.00	38.30
30	ATOM	291	CA	GLU	38	-54.617	115.797	-36.457	1.00	41.17
	ATOM	292	CB	GLU	38	-54.900	117.001	-37.348	1.00	46.15
	ATOM	293	CG	GLU	38	-53.900	118.129	-37.256	1.00	50.01
	ATOM	294	CD	GLU	38	-54.104	119.150	-38.369	1.00	51.18
	ATOM	295	OE1	GLU	38	-55.284	119.446	-38.689	1.00	48.40
35	ATOM	296	OE2	GLU	38	-53.086	119.650	-38.918	1.00	50.23
	ATOM	297	C	GLU	38	-55.949	115.171	-36.091	1.00	42.69
	ATOM	298	O	GLU	38	-56.380	114.200	-36.698	1.00	42.55
	ATOM	299	N	GLU	39	-56.608	115.734	-35.092	1.00	44.75
	ATOM	300	CA	GLU	39	-57.912	115.224	-34.702	1.00	46.50
40	ATOM	301	CB	GLU	39	-58.109	115.350	-33.194	1.00	47.52
	ATOM	302	CG	GLU	39	-57.533	116.631	-32.630	1.00	58.50
	ATOM	303	CD	GLU	39	-57.854	116.814	-31.159	1.00	62.35
	ATOM	304	OE1	GLU	39	-57.271	117.733	-30.528	1.00	64.58
	ATOM	305	OE2	GLU	39	-58.697	116.045	-30.639	1.00	63.04
45	ATOM	306	C	GLU	39	-58.891	116.117	-35.442	1.00	44.56
	ATOM	307	O	GLU	39	-58.696	117.323	-35.516	1.00	48.11
	ATOM	308	N	LYS	40	-59.930	115.528	-36.010	1.00	40.71
	ATOM	309	CA	LYS	40	-60.908	116.301	-36.749	1.00	38.08
	ATOM	310	CB	LYS	40	-60.469	116.454	-38.204	1.00	41.28
50	ATOM	311	CG	LYS	40	-61.516	117.098	-39.104	1.00	41.52
	ATOM	312	CD	LYS	40	-61.157	116.917	-40.572	1.00	42.73
	ATOM	313	CE	LYS	40	-62.258	117.406	-41.497	1.00	41.97
	ATOM	314	NZ	LYS	40	-61.930	117.102	-42.924	1.00	45.94
	ATOM	315	C	LYS	40	-62.271	115.643	-36.706	1.00	39.66
55	ATOM	316	O	LYS	40	-62.469	114.560	-37.251	1.00	36.18
	ATOM	317	N	GLU	41	-63.210	116.310	-36.047	1.00	40.16
	ATOM	318	CA	GLU	41	-64.575	115.821	-35.937	1.00	42.80
	ATOM	319	CB	GLU	41	-65.288	116.046	-37.265	1.00	45.31
	ATOM	320	CG	GLU	41	-65.218	117.503	-37.706	1.00	56.07
60	ATOM	321	CD	GLU	41	-65.524	117.704	-39.188	1.00	59.42
	ATOM	322	OE1	GLU	41	-64.864	117.059	-40.040	1.00	63.71
	ATOM	323	OE2	GLU	41	-66.424	118.515	-39.500	1.00	63.71
	ATOM	324	C	GLU	41	-64.638	114.358	-35.529	1.00	38.93
	ATOM	325	O	GLU	41	-65.266	113.542	-36.198	1.00	39.03
65	ATOM	326	N	ASN	42	-63.962	114.040	-34.434	1.00	34.69
	ATOM	327	CA	ASN	42	-63.940	112.695	-33.886	1.00	32.96

	ATOM	328	CB	ASN	42	-65.364	112.210	-33.638	1.00	32.65
	ATOM	329	CG	ASN	42	-65.462	111.338	-32.425	1.00	35.50
	ATOM	330	OD1	ASN	42	-66.119	110.307	-32.443	1.00	40.26
	ATOM	331	ND2	ASN	42	-64.808	111.754	-31.351	1.00	29.28
5	ATOM	332	C	ASN	42	-63.209	111.659	-34.726	1.00	31.44
	ATOM	333	O	ASN	42	-63.392	110.461	-34.537	1.00	22.78
	ATOM	334	N	LYS	43	-62.376	112.123	-35.644	1.00	33.26
	ATOM	335	CA	LYS	43	-61.620	111.228	-36.504	1.00	33.68
	ATOM	336	CB	LYS	43	-62.211	111.227	-37.914	1.00	33.55
10	ATOM	337	CG	LYS	43	-63.585	110.613	-37.999	1.00	40.49
	ATOM	338	CD	LYS	43	-64.145	110.731	-39.394	1.00	45.39
	ATOM	339	CE	LYS	43	-64.380	112.183	-39.757	1.00	52.57
	ATOM	340	NZ	LYS	43	-65.029	112.307	-41.088	1.00	62.06
	ATOM	341	C	LYS	43	-60.180	111.680	-36.566	1.00	32.05
15	ATOM	342	O	LYS	43	-59.859	112.789	-36.158	1.00	37.32
	ATOM	343	N	ILE	44	-59.303	110.820	-37.060	1.00	31.31
	ATOM	344	CA	ILE	44	-57.909	111.199	-37.172	1.00	29.52
	ATOM	345	CB	ILE	44	-56.982	110.058	-36.752	1.00	26.38
	ATOM	346	CG2	ILE	44	-55.538	110.481	-36.929	1.00	26.09
20	ATOM	347	CG1	ILE	44	-57.235	109.712	-35.284	1.00	28.61
	ATOM	348	CD1	ILE	44	-56.364	108.618	-34.746	1.00	21.51
	ATOM	349	C	ILE	44	-57.629	111.596	-38.611	1.00	31.60
	ATOM	350	O	ILE	44	-57.846	110.817	-39.536	1.00	34.50
	ATOM	351	N	LEU	45	-57.165	112.825	-38.792	1.00	31.44
25	ATOM	352	CA	LEU	45	-56.854	113.347	-40.114	1.00	31.40
	ATOM	353	CB	LEU	45	-57.314	114.799	-40.212	1.00	33.60
	ATOM	354	CG	LEU	45	-56.955	115.530	-41.503	1.00	34.36
	ATOM	355	CD1	LEU	45	-57.703	114.925	-42.673	1.00	29.00
	ATOM	356	CD2	LEU	45	-57.299	116.999	-41.347	1.00	32.42
30	ATOM	357	C	LEU	45	-55.365	113.261	-40.434	1.00	27.80
	ATOM	358	O	LEU	45	-54.522	113.691	-39.654	1.00	27.94
	ATOM	359	N	VAL	46	-55.054	112.697	-41.595	1.00	28.00
	ATOM	360	CA	VAL	46	-53.676	112.558	-42.044	1.00	31.53
	ATOM	361	CB	VAL	46	-53.547	111.344	-42.979	1.00	27.89
35	ATOM	362	CG1	VAL	46	-52.112	111.160	-43.410	1.00	26.10
	ATOM	363	CG2	VAL	46	-54.050	110.112	-42.278	1.00	23.39
	ATOM	364	C	VAL	46	-53.249	113.830	-42.792	1.00	33.94
	ATOM	365	O	VAL	46	-53.901	114.237	-43.756	1.00	39.68
	ATOM	366	N	LYS	47	-52.157	114.452	-42.353	1.00	33.59
40	ATOM	367	CA	LYS	47	-51.674	115.675	-42.990	1.00	37.05
	ATOM	368	CB	LYS	47	-51.407	116.752	-41.934	1.00	34.49
	ATOM	369	CG	LYS	47	-52.548	116.938	-40.949	1.00	37.53
	ATOM	370	CD	LYS	47	-53.853	117.279	-41.656	1.00	41.69
	ATOM	371	CE	LYS	47	-54.075	118.778	-41.713	1.00	42.95
45	ATOM	372	NZ	LYS	47	-52.856	119.493	-42.155	1.00	44.50
	ATOM	373	C	LYS	47	-50.420	115.465	-43.839	1.00	37.34
	ATOM	374	O	LYS	47	-50.026	116.345	-44.598	1.00	43.04
	ATOM	375	N	GLU	48	-49.797	114.303	-43.702	1.00	36.99
	ATOM	376	CA	GLU	48	-48.601	113.960	-44.466	1.00	37.31
50	ATOM	377	CB	GLU	48	-47.356	114.046	-43.599	1.00	36.53
	ATOM	378	CG	GLU	48	-46.960	115.429	-43.177	1.00	46.74
	ATOM	379	CD	GLU	48	-45.737	115.405	-42.278	1.00	52.39
	ATOM	380	OE1	GLU	48	-44.807	114.602	-42.564	1.00	51.33
	ATOM	381	OE2	GLU	48	-45.704	116.187	-41.293	1.00	54.55
55	ATOM	382	C	GLU	48	-48.757	112.526	-44.908	1.00	36.94
	ATOM	383	O	GLU	48	-49.027	111.665	-44.081	1.00	40.21
	ATOM	384	N	THR	49	-48.575	112.242	-46.191	1.00	36.34
	ATOM	385	CA	THR	49	-48.743	110.862	-46.619	1.00	35.48
	ATOM	386	CB	THR	49	-49.006	110.754	-48.147	1.00	31.16
60	ATOM	387	OG1	THR	49	-47.787	110.456	-48.824	1.00	32.48
	ATOM	388	CG2	THR	49	-49.569	112.046	-48.687	1.00	30.36
	ATOM	389	C	THR	49	-47.523	110.025	-46.225	1.00	34.59
	ATOM	390	O	THR	49	-46.409	110.534	-46.122	1.00	35.13
	ATOM	391	N	GLY	50	-47.757	108.737	-45.985	1.00	32.49
65	ATOM	392	CA	GLY	50	-46.683	107.846	-45.603	1.00	28.17
	ATOM	393	C	GLY	50	-47.229	106.569	-45.005	1.00	29.99

	ATOM	394	O	GLY	50	-48.389	106.244	-45.196	1.00	29.76
	ATOM	395	N	TYR	51	-46.382	105.840	-44.287	1.00	31.68
	ATOM	396	CA	TYR	51	-46.785	104.596	-43.639	1.00	32.88
	ATOM	397	CB	TYR	51	-45.715	103.515	-43.828	1.00	35.97
5	ATOM	398	CG	TYR	51	-45.631	103.018	-45.241	1.00	43.03
	ATOM	399	CD1	TYR	51	-44.924	103.731	-46.213	1.00	44.15
	ATOM	400	CE1	TYR	51	-44.920	103.322	-47.548	1.00	47.81
	ATOM	401	CD2	TYR	51	-46.329	101.878	-45.631	1.00	47.08
	ATOM	402	CE2	TYR	51	-46.339	101.458	-46.961	1.00	51.79
10	ATOM	403	CZ	TYR	51	-45.635	102.186	-47.918	1.00	51.15
	ATOM	404	OH	TYR	51	-45.689	101.798	-49.244	1.00	52.20
	ATOM	405	C	TYR	51	-47.037	104.817	-42.154	1.00	31.97
	ATOM	406	O	TYR	51	-46.186	105.339	-41.437	1.00	30.01
	ATOM	407	N	PHE	52	-48.216	104.417	-41.693	1.00	31.50
15	ATOM	408	CA	PHE	52	-48.568	104.592	-40.294	1.00	30.41
	ATOM	409	CB	PHE	52	-49.758	105.544	-40.156	1.00	31.30
	ATOM	410	CG	PHE	52	-49.525	106.910	-40.724	1.00	29.70
	ATOM	411	CD1	PHE	52	-49.502	107.113	-42.093	1.00	26.20
	ATOM	412	CD2	PHE	52	-49.340	107.997	-39.883	1.00	28.38
20	ATOM	413	CE1	PHE	52	-49.298	108.375	-42.613	1.00	27.17
	ATOM	414	CE2	PHE	52	-49.134	109.264	-40.396	1.00	27.76
	ATOM	415	CZ	PHE	52	-49.113	109.454	-41.761	1.00	27.27
	ATOM	416	C	PHE	52	-48.927	103.300	-39.592	1.00	27.65
	ATOM	417	O	PHE	52	-49.477	102.384	-40.190	1.00	24.85
25	ATOM	418	N	PHE	53	-48.598	103.242	-38.310	1.00	27.18
	ATOM	419	CA	PHE	53	-48.938	102.106	-37.468	1.00	27.05
	ATOM	420	CB	PHE	53	-47.859	101.861	-36.422	1.00	28.76
	ATOM	421	CG	PHE	53	-48.233	100.832	-35.404	1.00	28.00
	ATOM	422	CD1	PHE	53	-48.310	99.490	-35.745	1.00	26.49
30	ATOM	423	CD2	PHE	53	-48.528	101.206	-34.101	1.00	28.04
	ATOM	424	CE1	PHE	53	-48.674	98.536	-34.802	1.00	26.54
	ATOM	425	CE2	PHE	53	-48.895	100.253	-33.154	1.00	25.78
	ATOM	426	CZ	PHE	53	-48.966	98.919	-33.509	1.00	24.39
	ATOM	427	C	PHE	53	-50.215	102.585	-36.783	1.00	27.56
35	ATOM	428	O	PHE	53	-50.205	103.610	-36.108	1.00	25.39
	ATOM	429	N	ILE	54	-51.309	101.855	-36.974	1.00	28.87
	ATOM	430	CA	ILE	54	-52.596	102.238	-36.401	1.00	28.95
	ATOM	431	CB	ILE	54	-53.639	102.398	-37.526	1.00	29.60
	ATOM	432	CG2	ILE	54	-54.893	103.061	-36.981	1.00	27.42
40	ATOM	433	CG1	ILE	54	-53.046	103.247	-38.653	1.00	26.41
	ATOM	434	CD1	ILE	54	-53.818	103.208	-39.936	1.00	28.70
	ATOM	435	C	ILE	54	-53.084	101.207	-35.384	1.00	28.57
	ATOM	436	O	ILE	54	-52.993	100.005	-35.610	1.00	29.17
	ATOM	437	N	TYR	55	-53.611	101.676	-34.260	1.00	26.97
45	ATOM	438	CA	TYR	55	-54.072	100.758	-33.225	1.00	24.22
	ATOM	439	CB	TYR	55	-52.972	100.572	-32.180	1.00	21.82
	ATOM	440	CG	TYR	55	-52.530	101.849	-31.510	1.00	27.68
	ATOM	441	CD1	TYR	55	-53.142	102.294	-30.342	1.00	27.01
	ATOM	442	CE1	TYR	55	-52.751	103.477	-29.740	1.00	27.44
50	ATOM	443	CD2	TYR	55	-51.512	102.624	-32.054	1.00	27.16
	ATOM	444	CE2	TYR	55	-51.117	103.811	-31.458	1.00	25.39
	ATOM	445	CZ	TYR	55	-51.739	104.231	-30.305	1.00	29.02
	ATOM	446	OH	TYR	55	-51.354	105.412	-29.723	1.00	31.82
	ATOM	447	C	TYR	55	-55.353	101.208	-32.556	1.00	25.16
55	ATOM	448	O	TYR	55	-55.739	102.350	-32.661	1.00	28.47
	ATOM	449	N	GLY	56	-56.021	100.294	-31.873	1.00	27.44
	ATOM	450	CA	GLY	56	-57.255	100.644	-31.205	1.00	22.07
	ATOM	451	C	GLY	56	-57.731	99.553	-30.272	1.00	25.83
	ATOM	452	O	GLY	56	-57.526	98.372	-30.534	1.00	26.00
60	ATOM	453	N	GLN	57	-58.347	99.954	-29.167	1.00	22.56
	ATOM	454	CA	GLN	57	-58.882	99.017	-28.193	1.00	20.91
	ATOM	455	CB	GLN	57	-57.914	98.792	-27.030	1.00	17.87
	ATOM	456	CG	GLN	57	-58.510	97.918	-25.933	1.00	16.86
	ATOM	457	CD	GLN	57	-57.545	97.609	-24.808	1.00	22.06
65	ATOM	458	OE1	GLN	57	-56.533	96.946	-25.011	1.00	27.26
	ATOM	459	NE2	GLN	57	-57.859	98.084	-23.609	1.00	20.20

	ATOM	460	C	GLN	57	-60.187	99.562	-27.650	1.00	24.23
	ATOM	461	O	GLN	57	-60.338	100.772	-27.488	1.00	25.16
	ATOM	462	N	VAL	58	-61.122	98.658	-27.377	1.00	24.62
	ATOM	463	CA	VAL	58	-62.433	99.004	-26.841	1.00	26.68
5	ATOM	464	CB	VAL	58	-63.516	98.991	-27.947	1.00	25.55
	ATOM	465	CG1	VAL	58	-64.887	99.144	-27.335	1.00	15.69
	ATOM	466	CG2	VAL	58	-63.267	100.101	-28.945	1.00	25.77
	ATOM	467	C	VAL	58	-62.805	97.951	-25.818	1.00	31.06
	ATOM	468	O	VAL	58	-62.532	96.777	-26.033	1.00	27.74
10	ATOM	469	N	LEU	59	-63.417	98.363	-24.707	1.00	33.82
	ATOM	470	CA	LEU	59	-63.862	97.419	-23.676	1.00	29.85
	ATOM	471	CB	LEU	59	-63.629	97.974	-22.274	1.00	29.62
	ATOM	472	CG	LEU	59	-63.517	96.963	-21.125	1.00	28.11
	ATOM	473	CD1	LEU	59	-63.644	97.695	-19.804	1.00	26.67
15	ATOM	474	CD2	LEU	59	-64.572	95.901	-21.221	1.00	22.45
	ATOM	475	C	LEU	59	-65.359	97.189	-23.869	1.00	30.98
	ATOM	476	O	LEU	59	-66.162	98.104	-23.702	1.00	32.52
	ATOM	477	N	TYR	60	-65.729	95.964	-24.225	1.00	33.01
	ATOM	478	CA	TYR	60	-67.131	95.634	-24.439	1.00	34.51
20	ATOM	479	CB	TYR	60	-67.263	94.617	-25.568	1.00	40.10
	ATOM	480	CG	TYR	60	-66.757	95.143	-26.875	1.00	45.37
	ATOM	481	CD1	TYR	60	-65.542	94.708	-27.405	1.00	47.13
	ATOM	482	CE1	TYR	60	-65.037	95.261	-28.573	1.00	50.28
	ATOM	483	CD2	TYR	60	-67.455	96.136	-27.549	1.00	44.35
25	ATOM	484	CE2	TYR	60	-66.966	96.697	-28.712	1.00	48.81
	ATOM	485	CZ	TYR	60	-65.757	96.261	-29.216	1.00	51.91
	ATOM	486	OH	TYR	60	-65.260	96.857	-30.349	1.00	53.77
	ATOM	487	C	TYR	60	-67.818	95.101	-23.193	1.00	35.49
	ATOM	488	O	TYR	60	-67.395	94.111	-22.606	1.00	36.13
30	ATOM	489	N	THR	61	-68.887	95.776	-22.797	1.00	36.19
	ATOM	490	CA	THR	61	-69.660	95.378	-21.636	1.00	35.36
	ATOM	491	CB	THR	61	-69.651	96.477	-20.572	1.00	34.22
	ATOM	492	OG1	THR	61	-70.064	97.713	-21.162	1.00	34.22
	ATOM	493	CG2	THR	61	-68.260	96.635	-19.986	1.00	25.45
35	ATOM	494	C	THR	61	-71.082	95.131	-22.120	1.00	37.82
	ATOM	495	O	THR	61	-72.054	95.363	-21.407	1.00	43.70
	ATOM	496	N	ASP	62	-71.176	94.658	-23.355	1.00	37.64
	ATOM	497	CA	ASP	62	-72.433	94.362	-24.018	1.00	38.08
	ATOM	498	CB	ASP	62	-72.395	95.001	-25.402	1.00	38.42
40	ATOM	499	CG	ASP	62	-73.710	94.923	-26.123	1.00	41.25
	ATOM	500	OD1	ASP	62	-74.066	95.916	-26.801	1.00	38.28
	ATOM	501	OD2	ASP	62	-74.373	93.871	-26.019	1.00	41.37
	ATOM	502	C	ASP	62	-72.543	92.843	-24.120	1.00	40.90
	ATOM	503	O	ASP	62	-71.545	92.179	-24.365	1.00	46.73
45	ATOM	504	N	LYS	63	-73.734	92.280	-23.934	1.00	42.62
	ATOM	505	CA	LYS	63	-73.861	90.823	-24.004	1.00	40.92
	ATOM	506	CB	LYS	63	-74.755	90.313	-22.872	1.00	42.79
	ATOM	507	CG	LYS	63	-76.158	90.880	-22.861	1.00	45.91
	ATOM	508	CD	LYS	63	-76.884	90.476	-21.578	1.00	46.22
50	ATOM	509	CE	LYS	63	-76.168	91.039	-20.336	1.00	49.97
	ATOM	510	NZ	LYS	63	-76.691	90.502	-19.032	1.00	49.98
	ATOM	511	C	LYS	63	-74.359	90.271	-25.336	1.00	42.59
	ATOM	512	O	LYS	63	-74.819	89.139	-25.404	1.00	42.46
	ATOM	513	N	THR	64	-74.223	91.060	-26.394	1.00	44.06
55	ATOM	514	CA	THR	64	-74.663	90.665	-27.725	1.00	44.94
	ATOM	515	CB	THR	64	-74.651	91.888	-28.676	1.00	43.79
	ATOM	516	OG1	THR	64	-75.441	92.936	-28.107	1.00	46.20
	ATOM	517	CG2	THR	64	-75.234	91.531	-30.039	1.00	42.80
	ATOM	518	C	THR	64	-73.890	89.512	-28.386	1.00	45.82
60	ATOM	519	O	THR	64	-73.566	89.583	-29.557	1.00	53.10
	ATOM	520	N	TYR	65	-73.607	88.451	-27.647	1.00	45.11
	ATOM	521	CA	TYR	65	-72.909	87.268	-28.174	1.00	45.72
	ATOM	522	CB	TYR	65	-73.873	86.427	-29.032	1.00	40.46
	ATOM	523	CG	TYR	65	-73.845	86.700	-30.519	1.00	41.90
65	ATOM	524	CD1	TYR	65	-72.946	86.038	-31.357	1.00	42.47
	ATOM	525	CE1	TYR	65	-72.910	86.296	-32.739	1.00	41.83

	ATOM	526	CD2	TYR	65	-74.716	87.630	-31.096	1.00	41.65
	ATOM	527	CE2	TYR	65	-74.690	87.900	-32.475	1.00	40.28
	ATOM	528	CZ	TYR	65	-73.783	87.229	-33.286	1.00	41.90
	ATOM	529	OH	TYR	65	-73.753	87.488	-34.633	1.00	43.84
5	ATOM	530	C	TYR	65	-71.597	87.490	-28.944	1.00	45.31
	ATOM	531	O	TYR	65	-70.705	86.632	-28.923	1.00	49.74
	ATOM	532	N	ALA	66	-71.475	88.621	-29.625	1.00	42.32
	ATOM	533	CA	ALA	66	-70.273	88.929	-30.388	1.00	39.57
	ATOM	534	CB	ALA	66	-70.306	88.223	-31.725	1.00	35.36
10	ATOM	535	C	ALA	66	-70.164	90.429	-30.591	1.00	39.85
	ATOM	536	O	ALA	66	-71.062	91.059	-31.152	1.00	37.60
	ATOM	537	N	MET	67	-69.066	91.002	-30.115	1.00	41.88
	ATOM	538	CA	MET	67	-68.835	92.433	-30.250	1.00	41.48
	ATOM	539	CB	MET	67	-68.829	93.119	-28.879	1.00	39.98
15	ATOM	540	CG	MET	67	-70.180	93.157	-28.176	1.00	37.31
	ATOM	541	SD	MET	67	-71.445	94.011	-29.125	1.00	35.46
	ATOM	542	CE	MET	67	-71.077	95.729	-28.763	1.00	34.06
	ATOM	543	C	MET	67	-67.505	92.660	-30.946	1.00	39.50
	ATOM	544	O	MET	67	-66.694	91.740	-31.078	1.00	38.44
20	ATOM	545	N	GLY	68	-67.287	93.890	-31.394	1.00	39.84
	ATOM	546	CA	GLY	68	-66.048	94.216	-32.071	1.00	37.61
	ATOM	547	C	GLY	68	-66.091	95.584	-32.707	1.00	33.76
	ATOM	548	O	GLY	68	-67.134	96.228	-32.730	1.00	35.58
	ATOM	549	N	HIS	69	-64.953	96.040	-33.212	1.00	35.07
25	ATOM	550	CA	HIS	69	-64.883	97.340	-33.856	1.00	33.70
	ATOM	551	CB	HIS	69	-64.277	98.393	-32.909	1.00	37.02
	ATOM	552	CG	HIS	69	-62.952	98.012	-32.325	1.00	34.65
	ATOM	553	CD2	HIS	69	-61.701	98.446	-32.600	1.00	35.63
	ATOM	554	ND1	HIS	69	-62.819	97.086	-31.313	1.00	37.68
30	ATOM	555	CE1	HIS	69	-61.545	96.966	-30.989	1.00	32.64
	ATOM	556	NE2	HIS	69	-60.845	97.782	-31.755	1.00	35.90
	ATOM	557	C	HIS	69	-64.085	97.281	-35.151	1.00	33.98
	ATOM	558	O	HIS	69	-63.439	96.287	-35.444	1.00	30.94
	ATOM	559	N	LEU	70	-64.142	98.354	-35.927	1.00	35.35
35	ATOM	560	CA	LEU	70	-63.427	98.430	-37.191	1.00	30.76
	ATOM	561	CB	LEU	70	-64.411	98.505	-38.353	1.00	29.59
	ATOM	562	CG	LEU	70	-65.673	97.649	-38.314	1.00	30.71
	ATOM	563	CD1	LEU	70	-66.600	98.083	-39.423	1.00	25.60
	ATOM	564	CD2	LEU	70	-65.312	96.188	-38.441	1.00	27.39
40	ATOM	565	C	LEU	70	-62.572	99.692	-37.238	1.00	33.19
	ATOM	566	O	LEU	70	-63.008	100.760	-36.809	1.00	37.07
	ATOM	567	N	ILE	71	-61.346	99.573	-37.733	1.00	30.93
	ATOM	568	CA	ILE	71	-60.505	100.749	-37.890	1.00	27.94
	ATOM	569	CB	ILE	71	-59.043	100.473	-37.532	1.00	28.22
45	ATOM	570	CG2	ILE	71	-58.161	101.600	-38.030	1.00	24.54
	ATOM	571	CG1	ILE	71	-58.904	100.351	-36.019	1.00	26.34
	ATOM	572	CD1	ILE	71	-57.546	99.885	-35.573	1.00	37.50
	ATOM	573	C	ILE	71	-60.645	100.993	-39.383	1.00	30.32
	ATOM	574	O	ILE	71	-60.233	100.163	-40.185	1.00	23.07
50	ATOM	575	N	GLN	72	-61.254	102.115	-39.758	1.00	29.74
	ATOM	576	CA	GLN	72	-61.471	102.392	-41.171	1.00	27.85
	ATOM	577	CB	GLN	72	-62.960	102.577	-41.420	1.00	26.70
	ATOM	578	CG	GLN	72	-63.783	101.463	-40.840	1.00	28.28
	ATOM	579	CD	GLN	72	-65.224	101.540	-41.255	1.00	33.11
55	ATOM	580	OE1	GLN	72	-65.872	102.568	-41.078	1.00	41.22
	ATOM	581	NE2	GLN	72	-65.742	100.452	-41.807	1.00	30.18
	ATOM	582	C	GLN	72	-60.715	103.576	-41.735	1.00	29.85
	ATOM	583	O	GLN	72	-60.307	104.481	-41.004	1.00	28.20
	ATOM	584	N	ARG	73	-60.552	103.560	-43.052	1.00	30.90
60	ATOM	585	CA	ARG	73	-59.855	104.620	-43.763	1.00	30.79
	ATOM	586	CB	ARG	73	-58.594	104.056	-44.402	1.00	29.18
	ATOM	587	CG	ARG	73	-57.840	105.045	-45.239	1.00	28.66
	ATOM	588	CD	ARG	73	-56.989	104.328	-46.244	1.00	29.25
	ATOM	589	NE	ARG	73	-56.203	105.257	-47.041	1.00	35.78
65	ATOM	590	CZ	ARG	73	-55.645	104.953	-48.203	1.00	35.27
	ATOM	591	NH1	ARG	73	-55.793	103.739	-48.713	1.00	34.95

	ATOM	592	NH2	ARG	73	-54.938	105.862	-48.849	1.00	33.33
	ATOM	593	C	ARG	73	-60.728	105.245	-44.849	1.00	32.42
	ATOM	594	O	ARG	73	-61.292	104.537	-45.676	1.00	34.19
	ATOM	595	N	LYS	74	-60.849	106.569	-44.828	1.00	35.90
5	ATOM	596	CA	LYS	74	-61.613	107.289	-45.846	1.00	38.57
	ATOM	597	CB	LYS	74	-62.433	108.426	-45.230	1.00	43.52
	ATOM	598	CG	LYS	74	-63.647	107.978	-44.433	1.00	52.89
	ATOM	599	CD	LYS	74	-64.403	109.169	-43.800	1.00	58.14
	ATOM	600	CE	LYS	74	-65.609	108.685	-42.993	1.00	56.75
10	ATOM	601	NZ	LYS	74	-66.335	109.809	-42.331	1.00	63.98
	ATOM	602	C	LYS	74	-60.606	107.892	-46.818	1.00	40.11
	ATOM	603	O	LYS	74	-59.926	108.872	-46.486	1.00	37.55
	ATOM	604	N	LYS	75	-60.509	107.311	-48.009	1.00	34.48
	ATOM	605	CA	LYS	75	-59.574	107.789	-49.020	1.00	34.24
15	ATOM	606	CB	LYS	75	-59.535	106.811	-50.198	1.00	36.16
	ATOM	607	CG	LYS	75	-59.171	105.364	-49.872	1.00	35.99
	ATOM	608	CD	LYS	75	-59.196	104.528	-51.159	1.00	40.91
	ATOM	609	CE	LYS	75	-58.814	103.076	-50.907	1.00	46.16
	ATOM	610	NZ	LYS	75	-58.876	102.193	-52.124	1.00	46.03
20	ATOM	611	C	LYS	75	-59.969	109.173	-49.545	1.00	30.92
	ATOM	612	O	LYS	75	-61.139	109.439	-49.750	1.00	27.47
	ATOM	613	N	VAL	76	-58.986	110.043	-49.764	1.00	32.82
	ATOM	614	CA	VAL	76	-59.241	111.383	-50.300	1.00	31.51
	ATOM	615	CB	VAL	76	-58.049	112.340	-50.123	1.00	29.19
25	ATOM	616	CG1	VAL	76	-58.550	113.734	-49.876	1.00	27.09
	ATOM	617	CG2	VAL	76	-57.159	111.878	-49.025	1.00	34.94
	ATOM	618	C	VAL	76	-59.424	111.247	-51.798	1.00	33.40
	ATOM	619	O	VAL	76	-60.275	111.898	-52.395	1.00	32.92
	ATOM	620	N	HIS	77	-58.588	110.404	-52.395	1.00	34.88
30	ATOM	621	CA	HIS	77	-58.614	110.154	-53.825	1.00	34.59
	ATOM	622	CB	HIS	77	-57.195	110.232	-54.398	1.00	35.92
	ATOM	623	CG	HIS	77	-56.516	111.545	-54.155	1.00	34.17
	ATOM	624	CD2	HIS	77	-55.205	111.868	-54.080	1.00	31.44
	ATOM	625	ND1	HIS	77	-57.210	112.726	-53.989	1.00	35.05
35	ATOM	626	CE1	HIS	77	-56.359	113.719	-53.819	1.00	29.84
	ATOM	627	NE2	HIS	77	-55.136	113.226	-53.873	1.00	38.35
	ATOM	628	C	HIS	77	-59.217	108.783	-54.080	1.00	35.63
	ATOM	629	O	HIS	77	-58.922	107.829	-53.360	1.00	40.24
	ATOM	630	N	VAL	78	-60.042	108.677	-55.121	1.00	37.37
40	ATOM	631	CA	VAL	78	-60.706	107.416	-55.402	1.00	41.32
	ATOM	632	CB	VAL	78	-62.230	107.587	-55.252	1.00	43.10
	ATOM	633	CG1	VAL	78	-62.918	106.252	-55.400	1.00	45.01
	ATOM	634	CG2	VAL	78	-62.551	108.155	-53.881	1.00	44.98
	ATOM	635	C	VAL	78	-60.399	106.649	-56.694	1.00	43.02
45	ATOM	636	O	VAL	78	-59.832	105.560	-56.624	1.00	49.79
	ATOM	637	N	PHE	79	-60.762	107.161	-57.863	1.00	41.33
	ATOM	638	CA	PHE	79	-60.487	106.416	-59.116	1.00	45.00
	ATOM	639	CB	PHE	79	-59.089	105.766	-59.134	1.00	40.51
	ATOM	640	CG	PHE	79	-57.963	106.713	-58.889	1.00	39.52
50	ATOM	641	CD1	PHE	79	-57.306	106.721	-57.663	1.00	36.80
	ATOM	642	CD2	PHE	79	-57.552	107.593	-59.881	1.00	37.89
	ATOM	643	CE1	PHE	79	-56.256	107.592	-57.428	1.00	37.52
	ATOM	644	CE2	PHE	79	-56.501	108.467	-59.654	1.00	35.59
	ATOM	645	CZ	PHE	79	-55.851	108.468	-58.425	1.00	37.58
55	ATOM	646	C	PHE	79	-61.463	105.275	-59.404	1.00	43.50
	ATOM	647	O	PHE	79	-61.636	104.364	-58.592	1.00	37.85
	ATOM	648	N	GLY	80	-62.065	105.319	-60.587	1.00	46.65
	ATOM	649	CA	GLY	80	-62.986	104.284	-61.014	1.00	48.38
	ATOM	650	C	GLY	80	-64.038	103.831	-60.025	1.00	47.89
60	ATOM	651	O	GLY	80	-64.727	104.650	-59.411	1.00	51.15
	ATOM	652	N	ASP	81	-64.157	102.514	-59.870	1.00	45.07
	ATOM	653	CA	ASP	81	-65.148	101.939	-58.979	1.00	43.99
	ATOM	654	CB	ASP	81	-65.778	100.698	-59.626	1.00	45.66
	ATOM	655	CG	ASP	81	-64.810	99.530	-59.734	1.00	47.49
65	ATOM	656	OD1	ASP	81	-63.586	99.728	-59.563	1.00	51.90
	ATOM	657	OD2	ASP	81	-65.274	98.404	-60.002	1.00	48.08

	ATOM	658	C	ASP	81	-64.620	101.594	-57.598	1.00	42.29
	ATOM	659	O	ASP	81	-65.186	100.743	-56.912	1.00	39.94
	ATOM	660	N	GLU	82	-63.537	102.248	-57.189	1.00	39.91
	ATOM	661	CA	GLU	82	-62.975	102.012	-55.861	1.00	36.58
5	ATOM	662	CB	GLU	82	-61.702	102.830	-55.631	1.00	34.91
	ATOM	663	CG	GLU	82	-60.434	102.317	-56.234	1.00	37.82
	ATOM	664	CD	GLU	82	-59.218	102.825	-55.477	1.00	40.75
	ATOM	665	OE1	GLU	82	-59.184	104.022	-55.129	1.00	38.15
	ATOM	666	OE2	GLU	82	-58.291	102.028	-55.221	1.00	47.99
10	ATOM	667	C	GLU	82	-63.971	102.471	-54.806	1.00	36.58
	ATOM	668	O	GLU	82	-64.732	103.409	-55.028	1.00	38.40
	ATOM	669	N	LEU	83	-63.963	101.808	-53.659	1.00	33.83
	ATOM	670	CA	LEU	83	-64.811	102.205	-52.547	1.00	32.70
	ATOM	671	CB	LEU	83	-65.125	101.009	-51.648	1.00	30.78
15	ATOM	672	CG	LEU	83	-66.406	100.224	-51.920	1.00	29.26
	ATOM	673	CD1	LEU	83	-66.598	100.003	-53.400	1.00	29.60
	ATOM	674	CD2	LEU	83	-66.339	98.909	-51.187	1.00	28.65
	ATOM	675	C	LEU	83	-63.912	103.177	-51.813	1.00	33.88
	ATOM	676	O	LEU	83	-62.743	102.882	-51.595	1.00	38.46
20	ATOM	677	N	SER	84	-64.435	104.338	-51.444	1.00	34.94
	ATOM	678	CA	SER	84	-63.610	105.320	-50.758	1.00	39.77
	ATOM	679	CB	SER	84	-64.180	106.724	-50.952	1.00	38.92
	ATOM	680	OG	SER	84	-65.497	106.793	-50.463	1.00	43.11
	ATOM	681	C	SER	84	-63.436	105.029	-49.270	1.00	39.89
25	ATOM	682	O	SER	84	-62.607	105.641	-48.597	1.00	42.52
	ATOM	683	N	LEU	85	-64.217	104.088	-48.762	1.00	40.92
	ATOM	684	CA	LEU	85	-64.136	103.710	-47.364	1.00	34.66
	ATOM	685	CB	LEU	85	-65.525	103.763	-46.728	1.00	35.12
	ATOM	686	CG	LEU	85	-65.716	103.681	-45.206	1.00	36.92
30	ATOM	687	CD1	LEU	85	-65.349	102.305	-44.706	1.00	38.24
	ATOM	688	CD2	LEU	85	-64.878	104.751	-44.524	1.00	36.08
	ATOM	689	C	LEU	85	-63.579	102.300	-47.313	1.00	35.30
	ATOM	690	O	LEU	85	-64.195	101.355	-47.798	1.00	32.85
	ATOM	691	N	VAL	86	-62.391	102.167	-46.744	1.00	33.26
35	ATOM	692	CA	VAL	86	-61.745	100.869	-46.626	1.00	34.57
	ATOM	693	CB	VAL	86	-60.315	100.876	-47.239	1.00	35.39
	ATOM	694	CG1	VAL	86	-59.611	99.565	-46.946	1.00	31.44
	ATOM	695	CG2	VAL	86	-60.391	101.097	-48.737	1.00	38.08
	ATOM	696	C	VAL	86	-61.617	100.514	-45.164	1.00	32.85
40	ATOM	697	O	VAL	86	-61.205	101.340	-44.354	1.00	35.10
	ATOM	698	N	THR	87	-61.978	99.294	-44.809	1.00	33.41
	ATOM	699	CA	THR	87	-61.821	98.899	-43.430	1.00	33.56
	ATOM	700	CB	THR	87	-63.103	98.199	-42.880	1.00	32.70
	ATOM	701	OG1	THR	87	-62.741	97.087	-42.052	1.00	34.29
45	ATOM	702	CG2	THR	87	-64.002	97.764	-44.007	1.00	34.95
	ATOM	703	C	THR	87	-60.571	98.025	-43.273	1.00	30.57
	ATOM	704	O	THR	87	-60.471	96.941	-43.842	1.00	28.08
	ATOM	705	N	LEU	88	-59.597	98.561	-42.543	1.00	28.42
	ATOM	706	CA	LEU	88	-58.357	97.869	-42.247	1.00	31.31
50	ATOM	707	CB	LEU	88	-57.200	98.861	-42.066	1.00	28.21
	ATOM	708	CG	LEU	88	-56.718	99.980	-42.995	1.00	29.12
	ATOM	709	CD1	LEU	88	-57.406	99.928	-44.314	1.00	31.12
	ATOM	710	CD2	LEU	88	-56.947	101.311	-42.322	1.00	28.72
	ATOM	711	C	LEU	88	-58.579	97.184	-40.887	1.00	41.15
55	ATOM	712	O	LEU	88	-59.320	97.684	-40.036	1.00	52.30
	ATOM	713	N	PHE	89	-57.971	96.034	-40.667	1.00	39.23
	ATOM	714	CA	PHE	89	-58.091	95.404	-39.350	1.00	42.27
	ATOM	715	CB	PHE	89	-57.426	96.338	-38.335	1.00	35.63
	ATOM	716	CG	PHE	89	-56.146	96.936	-38.856	1.00	35.06
60	ATOM	717	CD1	PHE	89	-55.862	98.285	-38.669	1.00	31.56
	ATOM	718	CD2	PHE	89	-55.278	96.166	-39.650	1.00	30.96
	ATOM	719	CE1	PHE	89	-54.744	98.865	-39.278	1.00	33.76
	ATOM	720	CE2	PHE	89	-54.165	96.732	-40.261	1.00	27.38
	ATOM	721	CZ	PHE	89	-53.898	98.084	-40.079	1.00	30.29
65	ATOM	722	C	PHE	89	-59.476	94.922	-38.866	1.00	39.99
	ATOM	723	O	PHE	89	-60.061	94.023	-39.479	1.00	45.34

	ATOM	724	N	ARG	90	-60.008	95.464	-37.781	1.00	38.92
	ATOM	725	CA	ARG	90	-61.300	94.958	-37.282	1.00	41.66
	ATOM	726	CB	ARG	90	-62.281	94.705	-38.446	1.00	39.24
	ATOM	727	CG	ARG	90	-63.191	93.473	-38.274	1.00	29.41
5	ATOM	728	CD	ARG	90	-64.164	93.292	-39.455	1.00	31.13
	ATOM	729	NE	ARG	90	-64.830	91.987	-39.432	1.00	31.75
	ATOM	730	CZ	ARG	90	-64.652	91.021	-40.337	1.00	28.86
	ATOM	731	NH1	ARG	90	-63.836	91.189	-41.364	1.00	25.48
	ATOM	732	NH2	ARG	90	-65.274	89.862	-40.197	1.00	29.84
10	ATOM	733	C	ARG	90	-61.202	93.657	-36.463	1.00	39.23
	ATOM	734	O	ARG	90	-60.593	92.682	-36.901	1.00	31.33
	ATOM	735	N	CYS	91	-61.804	93.651	-35.271	1.00	40.12
	ATOM	736	CA	CYS	91	-61.810	92.454	-34.445	1.00	37.09
	ATOM	737	C	CYS	91	-63.115	92.093	-33.794	1.00	37.21
15	ATOM	738	O	CYS	91	-64.045	92.889	-33.757	1.00	37.46
	ATOM	739	CB	CYS	91	-60.742	92.488	-33.377	1.00	40.13
	ATOM	740	SG	CYS	91	-60.698	93.759	-32.064	1.00	41.64
	ATOM	741	N	ILE	92	-63.162	90.872	-33.274	1.00	34.02
	ATOM	742	CA	ILE	92	-64.363	90.343	-32.644	1.00	33.22
20	ATOM	743	CB	ILE	92	-65.064	89.348	-33.591	1.00	32.88
	ATOM	744	CG2	ILE	92	-66.408	88.944	-33.024	1.00	32.61
	ATOM	745	CG1	ILE	92	-65.254	89.992	-34.966	1.00	34.64
	ATOM	746	CD1	ILE	92	-65.757	89.042	-36.041	1.00	32.30
	ATOM	747	C	ILE	92	-64.078	89.629	-31.329	1.00	30.78
25	ATOM	748	O	ILE	92	-62.996	89.122	-31.119	1.00	33.43
	ATOM	749	N	GLN	93	-65.059	89.620	-30.434	1.00	29.35
	ATOM	750	CA	GLN	93	-64.943	88.941	-29.154	1.00	28.69
	ATOM	751	CB	GLN	93	-64.564	89.904	-28.028	1.00	28.17
	ATOM	752	CG	GLN	93	-63.077	90.215	-27.842	1.00	25.87
30	ATOM	753	CD	GLN	93	-62.232	89.001	-27.525	1.00	24.22
	ATOM	754	OE1	GLN	93	-61.741	88.331	-28.421	1.00	28.90
	ATOM	755	NE2	GLN	93	-62.060	88.711	-26.242	1.00	25.29
	ATOM	756	C	GLN	93	-66.301	88.361	-28.831	1.00	31.44
	ATOM	757	O	GLN	93	-67.293	89.074	-28.853	1.00	32.07
35	ATOM	758	N	ASN	94	-66.365	87.064	-28.563	1.00	35.77
	ATOM	759	CA	ASN	94	-67.631	86.464	-28.190	1.00	33.56
	ATOM	760	CB	ASN	94	-67.531	84.938	-28.168	1.00	34.70
	ATOM	761	CG	ASN	94	-67.670	84.325	-29.539	1.00	33.01
	ATOM	762	OD1	ASN	94	-68.614	84.610	-30.257	1.00	30.56
40	ATOM	763	ND2	ASN	94	-66.740	83.464	-29.900	1.00	36.61
	ATOM	764	C	ASN	94	-67.922	86.994	-26.786	1.00	35.80
	ATOM	765	O	ASN	94	-67.008	87.168	-25.977	1.00	36.63
	ATOM	766	N	MET	95	-69.188	87.262	-26.496	1.00	35.73
	ATOM	767	CA	MET	95	-69.561	87.785	-25.187	1.00	34.73
45	ATOM	768	CB	MET	95	-70.387	89.063	-25.360	1.00	32.97
	ATOM	769	CG	MET	95	-69.733	90.142	-26.211	1.00	27.35
	ATOM	770	SD	MET	95	-68.087	90.622	-25.650	1.00	25.14
	ATOM	771	CE	MET	95	-68.427	91.541	-24.187	1.00	15.55
	ATOM	772	C	MET	95	-70.351	86.774	-24.354	1.00	36.36
50	ATOM	773	O	MET	95	-71.029	85.901	-24.903	1.00	40.70
	ATOM	774	N	PRO	96	-70.247	86.860	-23.017	1.00	36.06
	ATOM	775	CD	PRO	96	-69.201	87.581	-22.292	1.00	35.32
	ATOM	776	CA	PRO	96	-70.958	85.967	-22.101	1.00	40.00
	ATOM	777	CB	PRO	96	-70.105	85.996	-20.837	1.00	33.97
55	ATOM	778	CG	PRO	96	-68.814	86.563	-21.275	1.00	37.05
	ATOM	779	C	PRO	96	-72.328	86.590	-21.844	1.00	46.34
	ATOM	780	O	PRO	96	-72.556	87.749	-22.187	1.00	47.59
	ATOM	781	N	GLU	97	-73.235	85.838	-21.229	1.00	52.14
	ATOM	782	CA	GLU	97	-74.568	86.357	-20.946	1.00	57.27
60	ATOM	783	CB	GLU	97	-75.578	85.213	-20.854	1.00	62.29
	ATOM	784	CG	GLU	97	-77.024	85.622	-21.118	1.00	72.31
	ATOM	785	CD	GLU	97	-77.470	85.290	-22.550	1.00	79.34
	ATOM	786	OE1	GLU	97	-77.477	84.073	-22.915	1.00	84.94
	ATOM	787	OE2	GLU	97	-77.807	86.242	-23.308	1.00	81.39
65	ATOM	788	C	GLU	97	-74.540	87.105	-19.617	1.00	56.67
	ATOM	789	O	GLU	97	-75.214	88.125	-19.448	1.00	59.18

	ATOM	790	N	THR	98	-73.740	86.602	-18.685	1.00	57.55
	ATOM	791	CA	THR	98	-73.641	87.182	-17.350	1.00	59.71
	ATOM	792	CB	THR	98	-72.871	86.252	-16.422	1.00	59.24
	ATOM	793	OG1	THR	98	-71.559	86.035	-16.977	1.00	68.81
5	ATOM	794	CG2	THR	98	-73.625	84.915	-16.263	1.00	56.15
	ATOM	795	C	THR	98	-73.015	88.570	-17.237	1.00	60.32
	ATOM	796	O	THR	98	-73.710	89.579	-17.408	1.00	65.03
	ATOM	797	N	LEU	99	-71.718	88.628	-16.929	1.00	54.94
	ATOM	798	CA	LEU	99	-71.026	89.909	-16.763	1.00	51.13
10	ATOM	799	CB	LEU	99	-70.252	89.905	-15.447	1.00	50.56
	ATOM	800	CG	LEU	99	-71.078	90.015	-14.168	1.00	52.80
	ATOM	801	CD1	LEU	99	-70.252	89.569	-12.963	1.00	48.38
	ATOM	802	CD2	LEU	99	-71.552	91.452	-14.007	1.00	50.80
	ATOM	803	C	LEU	99	-70.077	90.260	-17.913	1.00	50.20
15	ATOM	804	O	LEU	99	-68.850	90.131	-17.784	1.00	50.50
	ATOM	805	N	PRO	100	-70.628	90.745	-19.040	1.00	47.97
	ATOM	806	CD	PRO	100	-72.032	91.130	-19.248	1.00	45.68
	ATOM	807	CA	PRO	100	-69.821	91.106	-20.209	1.00	46.54
	ATOM	808	CB	PRO	100	-70.840	91.742	-21.156	1.00	45.50
20	ATOM	809	CG	PRO	100	-72.133	91.116	-20.746	1.00	45.80
	ATOM	810	C	PRO	100	-68.689	92.063	-19.875	1.00	46.78
	ATOM	811	O	PRO	100	-68.915	93.143	-19.341	1.00	52.13
	ATOM	812	N	ASN	101	-67.470	91.660	-20.191	1.00	42.71
	ATOM	813	CA	ASN	101	-66.301	92.485	-19.957	1.00	41.37
25	ATOM	814	CB	ASN	101	-65.946	92.499	-18.475	1.00	43.24
	ATOM	815	CG	ASN	101	-66.742	93.525	-17.693	1.00	45.93
	ATOM	816	OD1	ASN	101	-66.546	94.732	-17.849	1.00	45.33
	ATOM	817	ND2	ASN	101	-67.655	93.049	-16.848	1.00	48.11
	ATOM	818	C	ASN	101	-65.152	91.911	-20.762	1.00	42.66
30	ATOM	819	O	ASN	101	-64.326	91.162	-20.238	1.00	44.79
	ATOM	820	N	ASN	102	-65.093	92.248	-22.047	1.00	40.46
	ATOM	821	CA	ASN	102	-64.019	91.720	-22.854	1.00	37.90
	ATOM	822	CB	ASN	102	-64.574	90.909	-24.015	1.00	34.01
	ATOM	823	CG	ASN	102	-64.827	89.464	-23.630	1.00	32.44
35	ATOM	824	OD1	ASN	102	-64.033	88.859	-22.915	1.00	29.07
	ATOM	825	ND2	ASN	102	-65.929	88.904	-24.106	1.00	33.79
	ATOM	826	C	ASN	102	-62.935	92.657	-23.349	1.00	39.20
	ATOM	827	O	ASN	102	-61.783	92.508	-22.959	1.00	46.07
	ATOM	828	N	SER	103	-63.255	93.624	-24.187	1.00	35.42
40	ATOM	829	CA	SER	103	-62.184	94.501	-24.691	1.00	40.28
	ATOM	830	CB	SER	103	-61.355	95.119	-23.538	1.00	40.68
	ATOM	831	OG	SER	103	-60.101	94.473	-23.351	1.00	31.76
	ATOM	832	C	SER	103	-61.250	93.725	-25.649	1.00	35.24
	ATOM	833	O	SER	103	-60.661	92.704	-25.300	1.00	24.65
45	ATOM	834	N	CYS	104	-61.157	94.222	-26.876	1.00	32.88
	ATOM	835	CA	CYS	104	-60.333	93.612	-27.888	1.00	31.77
	ATOM	836	C	CYS	104	-59.384	94.689	-28.443	1.00	30.58
	ATOM	837	O	CYS	104	-59.791	95.818	-28.700	1.00	29.23
	ATOM	838	CB	CYS	104	-61.214	93.005	-29.015	1.00	28.43
50	ATOM	839	SG	CYS	104	-60.144	92.616	-30.409	1.00	46.07
	ATOM	840	N	TYR	105	-58.108	94.339	-28.592	1.00	30.34
	ATOM	841	CA	TYR	105	-57.102	95.243	-29.145	1.00	26.12
	ATOM	842	CB	TYR	105	-55.869	95.294	-28.244	1.00	27.03
	ATOM	843	CG	TYR	105	-54.686	96.054	-28.822	1.00	25.61
55	ATOM	844	CD1	TYR	105	-54.436	97.385	-28.473	1.00	24.90
	ATOM	845	CE1	TYR	105	-53.357	98.079	-29.001	1.00	26.08
	ATOM	846	CD2	TYR	105	-53.816	95.445	-29.721	1.00	25.41
	ATOM	847	CE2	TYR	105	-52.736	96.135	-30.256	1.00	28.91
	ATOM	848	CZ	TYR	105	-52.512	97.451	-29.892	1.00	27.58
60	ATOM	849	OH	TYR	105	-51.433	98.127	-30.414	1.00	29.89
	ATOM	850	C	TYR	105	-56.690	94.716	-30.517	1.00	29.20
	ATOM	851	O	TYR	105	-56.616	93.510	-30.732	1.00	33.10
	ATOM	852	N	SER	106	-56.422	95.614	-31.451	1.00	30.48
	ATOM	853	CA	SER	106	-56.011	95.194	-32.780	1.00	29.22
65	ATOM	854	CB	SER	106	-57.240	94.847	-33.625	1.00	28.30
	ATOM	855	OG	SER	106	-56.876	94.384	-34.910	1.00	28.62

	ATOM	856	C	SER	106	-55.233	96.334	-33.414	1.00	29.94
	ATOM	857	O	SER	106	-55.566	97.494	-33.201	1.00	29.72
	ATOM	858	N	ALA	107	-54.190	96.006	-34.171	1.00	27.64
	ATOM	859	CA	ALA	107	-53.376	97.020	-34.828	1.00	24.33
5	ATOM	860	CB	ALA	107	-52.322	97.548	-33.870	1.00	21.30
	ATOM	861	C	ALA	107	-52.709	96.483	-36.085	1.00	25.56
	ATOM	862	O	ALA	107	-52.656	95.281	-36.312	1.00	28.03
	ATOM	863	N	GLY	108	-52.197	97.389	-36.902	1.00	26.07
	ATOM	864	CA	GLY	108	-51.526	96.989	-38.121	1.00	25.49
10	ATOM	865	C	GLY	108	-50.954	98.197	-38.826	1.00	29.03
	ATOM	866	O	GLY	108	-51.156	99.319	-38.383	1.00	32.20
	ATOM	867	N	ILE	109	-50.242	97.975	-39.921	1.00	29.34
	ATOM	868	CA	ILE	109	-49.645	99.065	-40.678	1.00	27.71
	ATOM	869	CB	ILE	109	-48.173	98.766	-41.003	1.00	27.48
15	ATOM	870	CG2	ILE	109	-47.572	99.899	-41.809	1.00	26.46
	ATOM	871	CG1	ILE	109	-47.395	98.567	-39.703	1.00	26.16
	ATOM	872	CD1	ILE	109	-45.980	98.105	-39.901	1.00	29.07
	ATOM	873	C	ILE	109	-50.408	99.267	-41.973	1.00	28.66
	ATOM	874	O	ILE	109	-50.857	98.307	-42.594	1.00	30.59
20	ATOM	875	N	ALA	110	-50.561	100.520	-42.380	1.00	25.60
	ATOM	876	CA	ALA	110	-51.270	100.841	-43.613	1.00	26.36
	ATOM	877	CB	ALA	110	-52.736	101.063	-43.325	1.00	20.26
	ATOM	878	C	ALA	110	-50.678	102.091	-44.239	1.00	29.61
	ATOM	879	O	ALA	110	-50.092	102.914	-43.552	1.00	27.61
25	ATOM	880	N	LYS	111	-50.808	102.229	-45.551	1.00	33.77
	ATOM	881	CA	LYS	111	-50.296	103.416	-46.195	1.00	35.09
	ATOM	882	CB	LYS	111	-49.711	103.103	-47.567	1.00	39.38
	ATOM	883	CG	LYS	111	-49.327	104.382	-48.300	1.00	49.74
	ATOM	884	CD	LYS	111	-48.106	104.225	-49.164	1.00	53.95
30	ATOM	885	CE	LYS	111	-47.686	105.578	-49.721	1.00	54.28
	ATOM	886	NZ	LYS	111	-46.483	105.438	-50.611	1.00	60.85
	ATOM	887	C	LYS	111	-51.442	104.412	-46.328	1.00	35.28
	ATOM	888	O	LYS	111	-52.498	104.087	-46.857	1.00	36.58
	ATOM	889	N	LEU	112	-51.221	105.621	-45.831	1.00	32.32
35	ATOM	890	CA	LEU	112	-52.228	106.665	-45.868	1.00	29.67
	ATOM	891	CB	LEU	112	-52.586	107.089	-44.446	1.00	27.04
	ATOM	892	CG	LEU	112	-52.919	105.963	-43.468	1.00	24.38
	ATOM	893	CD1	LEU	112	-53.114	106.531	-42.078	1.00	27.56
	ATOM	894	CD2	LEU	112	-54.153	105.236	-43.928	1.00	19.52
40	ATOM	895	C	LEU	112	-51.708	107.868	-46.640	1.00	32.58
	ATOM	896	O	LEU	112	-50.509	107.984	-46.891	1.00	31.51
	ATOM	897	N	GLU	113	-52.613	108.766	-47.010	1.00	35.75
	ATOM	898	CA	GLU	113	-52.241	109.956	-47.754	1.00	40.08
	ATOM	899	CB	GLU	113	-52.710	109.863	-49.194	1.00	44.23
45	ATOM	900	CG	GLU	113	-52.416	108.571	-49.896	1.00	50.12
	ATOM	901	CD	GLU	113	-52.440	108.761	-51.395	1.00	58.08
	ATOM	902	OE1	GLU	113	-53.311	109.537	-51.886	1.00	61.13
	ATOM	903	OE2	GLU	113	-51.586	108.137	-52.077	1.00	63.64
	ATOM	904	C	GLU	113	-52.856	111.203	-47.170	1.00	41.63
50	ATOM	905	O	GLU	113	-53.837	111.136	-46.442	1.00	48.29
	ATOM	906	N	GLU	114	-52.270	112.345	-47.515	1.00	40.97
	ATOM	907	CA	GLU	114	-52.761	113.645	-47.078	1.00	36.68
	ATOM	908	CB	GLU	114	-52.133	114.746	-47.906	1.00	41.19
	ATOM	909	CG	GLU	114	-50.942	115.420	-47.326	1.00	46.58
55	ATOM	910	CD	GLU	114	-50.816	116.825	-47.873	1.00	51.12
	ATOM	911	OE1	GLU	114	-51.702	117.657	-47.560	1.00	51.88
	ATOM	912	OE2	GLU	114	-49.846	117.090	-48.625	1.00	57.08
	ATOM	913	C	GLU	114	-54.251	113.714	-47.341	1.00	35.51
	ATOM	914	O	GLU	114	-54.684	113.549	-48.475	1.00	38.92
60	ATOM	915	N	GLY	115	-55.036	113.974	-46.308	1.00	32.99
	ATOM	916	CA	GLY	115	-56.463	114.077	-46.520	1.00	31.79
	ATOM	917	C	GLY	115	-57.235	112.852	-46.111	1.00	34.39
	ATOM	918	O	GLY	115	-58.446	112.921	-45.956	1.00	39.20
	ATOM	919	N	ASP	116	-56.550	111.726	-45.962	1.00	35.66
65	ATOM	920	CA	ASP	116	-57.220	110.508	-45.538	1.00	38.09
	ATOM	921	CB	ASP	116	-56.275	109.302	-45.595	1.00	36.00

	ATOM	922	CG	ASP	116	-56.062	108.782	-46.999	1.00	35.09
	ATOM	923	OD1	ASP	116	-56.844	109.137	-47.906	1.00	33.81
	ATOM	924	OD2	ASP	116	-55.117	107.996	-47.193	1.00	35.24
	ATOM	925	C	ASP	116	-57.666	110.714	-44.098	1.00	39.38
5	ATOM	926	O	ASP	116	-57.066	111.498	-43.357	1.00	37.91
	ATOM	927	N	GLU	117	-58.727	110.023	-43.702	1.00	41.54
	ATOM	928	CA	GLU	117	-59.208	110.118	-42.330	1.00	37.92
	ATOM	929	CB	GLU	117	-60.568	110.804	-42.274	1.00	38.90
	ATOM	930	CG	GLU	117	-60.579	112.192	-42.862	1.00	48.15
10	ATOM	931	CD	GLU	117	-61.939	112.848	-42.746	1.00	52.62
	ATOM	932	OE1	GLU	117	-62.968	112.135	-42.903	1.00	55.49
	ATOM	933	OE2	GLU	117	-61.974	114.080	-42.507	1.00	57.10
	ATOM	934	C	GLU	117	-59.332	108.712	-41.778	1.00	34.02
	ATOM	935	O	GLU	117	-59.673	107.784	-42.509	1.00	33.31
15	ATOM	936	N	LEU	118	-59.026	108.552	-40.498	1.00	30.68
	ATOM	937	CA	LEU	118	-59.137	107.261	-39.841	1.00	27.14
	ATOM	938	CB	LEU	118	-57.839	106.916	-39.110	1.00	26.49
	ATOM	939	CG	LEU	118	-56.580	106.683	-39.938	1.00	26.21
	ATOM	940	CD1	LEU	118	-55.410	106.445	-39.012	1.00	23.05
20	ATOM	941	CD2	LEU	118	-56.779	105.498	-40.860	1.00	25.38
	ATOM	942	C	LEU	118	-60.269	107.337	-38.825	1.00	28.33
	ATOM	943	O	LEU	118	-60.489	108.383	-38.213	1.00	31.46
	ATOM	944	N	GLN	119	-60.999	106.240	-38.650	1.00	28.96
	ATOM	945	CA	GLN	119	-62.074	106.216	-37.666	1.00	29.13
25	ATOM	946	CB	GLN	119	-63.372	106.709	-38.289	1.00	28.23
	ATOM	947	CG	GLN	119	-63.964	105.769	-39.295	1.00	36.38
	ATOM	948	CD	GLN	119	-65.212	106.329	-39.934	1.00	36.57
	ATOM	949	OE1	GLN	119	-65.978	105.594	-40.557	1.00	40.70
	ATOM	950	NE2	GLN	119	-65.421	107.632	-39.795	1.00	36.80
30	ATOM	951	C	GLN	119	-62.269	104.827	-37.066	1.00	29.43
	ATOM	952	O	GLN	119	-61.945	103.819	-37.689	1.00	28.80
	ATOM	953	N	LEU	120	-62.783	104.799	-35.842	1.00	32.26
	ATOM	954	CA	LEU	120	-63.042	103.569	-35.099	1.00	29.96
	ATOM	955	CB	LEU	120	-62.469	103.698	-33.680	1.00	31.17
35	ATOM	956	CG	LEU	120	-62.300	102.541	-32.688	1.00	30.26
	ATOM	957	CD1	LEU	120	-63.547	101.696	-32.615	1.00	27.10
	ATOM	958	CD2	LEU	120	-61.117	101.714	-33.106	1.00	30.96
	ATOM	959	C	LEU	120	-64.562	103.406	-35.036	1.00	30.51
	ATOM	960	O	LEU	120	-65.258	104.202	-34.404	1.00	32.88
40	ATOM	961	N	ALA	121	-65.074	102.369	-35.689	1.00	27.40
	ATOM	962	CA	ALA	121	-66.510	102.126	-35.719	1.00	25.55
	ATOM	963	CB	ALA	121	-67.010	102.240	-37.151	1.00	19.41
	ATOM	964	C	ALA	121	-66.935	100.779	-35.133	1.00	29.89
	ATOM	965	O	ALA	121	-66.289	99.757	-35.357	1.00	35.88
45	ATOM	966	N	ILE	122	-68.025	100.795	-34.371	1.00	30.33
	ATOM	967	CA	ILE	122	-68.569	99.581	-33.777	1.00	30.27
	ATOM	968	CB	ILE	122	-68.893	99.776	-32.292	1.00	29.27
	ATOM	969	CG2	ILE	122	-69.365	98.466	-31.705	1.00	30.38
	ATOM	970	CG1	ILE	122	-67.650	100.272	-31.548	1.00	26.03
50	ATOM	971	CD1	ILE	122	-67.859	100.534	-30.068	1.00	29.21
	ATOM	972	C	ILE	122	-69.856	99.285	-34.538	1.00	33.20
	ATOM	973	O	ILE	122	-70.811	100.057	-34.475	1.00	33.30
	ATOM	974	N	PRO	123	-69.894	98.162	-35.277	1.00	34.70
	ATOM	975	CD	PRO	123	-68.787	97.214	-35.438	1.00	34.40
55	ATOM	976	CA	PRO	123	-71.048	97.732	-36.078	1.00	36.49
	ATOM	977	CB	PRO	123	-70.473	96.622	-36.963	1.00	30.22
	ATOM	978	CG	PRO	123	-68.984	96.784	-36.854	1.00	36.72
	ATOM	979	C	PRO	123	-72.227	97.216	-35.254	1.00	39.64
	ATOM	980	O	PRO	123	-72.670	96.081	-35.450	1.00	40.48
60	ATOM	981	N	ARG	124	-72.731	98.044	-34.343	1.00	44.58
	ATOM	982	CA	ARG	124	-73.854	97.660	-33.500	1.00	48.04
	ATOM	983	CB	ARG	124	-73.356	96.976	-32.232	1.00	52.91
	ATOM	984	CG	ARG	124	-72.448	95.789	-32.493	1.00	60.50
	ATOM	985	CD	ARG	124	-73.189	94.517	-32.140	1.00	72.57
65	ATOM	986	NE	ARG	124	-72.431	93.312	-32.521	1.00	84.90
	ATOM	987	CZ	ARG	124	-72.529	92.674	-33.694	1.00	85.44

	ATOM	988	NH1	ARG	124	-73.363	93.123	-34.634	1.00	87.69
	ATOM	989	NH2	ARG	124	-71.785	91.586	-33.929	1.00	82.78
	ATOM	990	C	ARG	124	-74.719	98.855	-33.134	1.00	48.39
	ATOM	991	O	ARG	124	-74.218	99.971	-32.991	1.00	45.26
5	ATOM	992	N	GLU	125	-76.015	98.595	-32.979	1.00	52.84
	ATOM	993	CA	GLU	125	-77.009	99.615	-32.652	1.00	55.02
	ATOM	994	CB	GLU	125	-78.358	98.944	-32.374	1.00	63.62
	ATOM	995	CG	GLU	125	-78.245	97.528	-31.806	1.00	70.62
	ATOM	996	CD	GLU	125	-77.404	96.609	-32.686	1.00	74.76
10	ATOM	997	OE1	GLU	125	-77.829	96.327	-33.839	1.00	75.75
	ATOM	998	OE2	GLU	125	-76.321	96.173	-32.223	1.00	80.71
	ATOM	999	C	GLU	125	-76.642	100.535	-31.496	1.00	53.44
	ATOM	1000	O	GLU	125	-76.377	101.719	-31.710	1.00	57.50
	ATOM	1001	N	ASN	126	-76.665	100.026	-30.273	1.00	47.18
15	ATOM	1002	CA	ASN	126	-76.286	100.857	-29.138	1.00	47.59
	ATOM	1003	CB	ASN	126	-77.494	101.243	-28.289	1.00	51.03
	ATOM	1004	CG	ASN	126	-77.962	102.665	-28.563	1.00	57.07
	ATOM	1005	OD1	ASN	126	-78.599	102.947	-29.590	1.00	61.15
	ATOM	1006	ND2	ASN	126	-77.629	103.578	-27.650	1.00	56.63
20	ATOM	1007	C	ASN	126	-75.295	100.082	-28.314	1.00	47.06
	ATOM	1008	O	ASN	126	-75.586	99.631	-27.203	1.00	45.61
	ATOM	1009	N	ALA	127	-74.111	99.927	-28.893	1.00	43.50
	ATOM	1010	CA	ALA	127	-73.038	99.184	-28.272	1.00	38.26
	ATOM	1011	CB	ALA	127	-71.755	99.423	-29.034	1.00	37.97
25	ATOM	1012	C	ALA	127	-72.850	99.547	-26.813	1.00	37.16
	ATOM	1013	O	ALA	127	-72.736	100.719	-26.460	1.00	33.14
	ATOM	1014	N	GLN	128	-72.844	98.527	-25.963	1.00	37.87
	ATOM	1015	CA	GLN	128	-72.610	98.734	-24.541	1.00	37.19
	ATOM	1016	CB	GLN	128	-73.291	97.631	-23.738	1.00	40.23
30	ATOM	1017	CG	GLN	128	-74.797	97.733	-23.801	1.00	38.14
	ATOM	1018	CD	GLN	128	-75.270	99.104	-23.379	1.00	39.51
	ATOM	1019	OE1	GLN	128	-75.151	99.486	-22.210	1.00	40.36
	ATOM	1020	NE2	GLN	128	-75.791	99.871	-24.337	1.00	41.53
	ATOM	1021	C	GLN	128	-71.091	98.691	-24.392	1.00	35.55
35	ATOM	1022	O	GLN	128	-70.468	97.626	-24.362	1.00	35.57
	ATOM	1023	N	ILE	129	-70.513	99.880	-24.312	1.00	34.64
	ATOM	1024	CA	ILE	129	-69.071	100.070	-24.256	1.00	29.60
	ATOM	1025	CB	ILE	129	-68.645	100.820	-25.571	1.00	31.03
	ATOM	1026	CG2	ILE	129	-67.720	101.978	-25.297	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-68.054	99.826	-26.545	1.00	26.37
	ATOM	1028	CD1	ILE	129	-69.017	98.769	-26.938	1.00	34.70
	ATOM	1029	C	ILE	129	-68.617	100.844	-23.022	1.00	29.55
	ATOM	1030	O	ILE	129	-69.378	101.611	-22.451	1.00	33.42
	ATOM	1031	N	SER	130	-67.375	100.633	-22.601	1.00	27.67
45	ATOM	1032	CA	SER	130	-66.836	101.373	-21.465	1.00	25.00
	ATOM	1033	CB	SER	130	-65.835	100.534	-20.688	1.00	22.09
	ATOM	1034	OG	SER	130	-65.155	101.347	-19.740	1.00	20.89
	ATOM	1035	C	SER	130	-66.121	102.599	-22.009	1.00	26.03
	ATOM	1036	O	SER	130	-65.394	102.498	-22.981	1.00	30.77
50	ATOM	1037	N	LEU	131	-66.317	103.753	-21.388	1.00	24.49
	ATOM	1038	CA	LEU	131	-65.678	104.969	-21.865	1.00	28.07
	ATOM	1039	CB	LEU	131	-66.714	106.082	-22.026	1.00	28.87
	ATOM	1040	CG	LEU	131	-67.427	106.160	-23.375	1.00	30.76
	ATOM	1041	CD1	LEU	131	-67.920	104.810	-23.806	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-68.574	107.129	-23.261	1.00	33.37
	ATOM	1043	C	LEU	131	-64.529	105.465	-21.004	1.00	33.66
	ATOM	1044	O	LEU	131	-64.280	106.669	-20.929	1.00	37.67
	ATOM	1045	N	ASP	132	-63.830	104.541	-20.354	1.00	34.97
	ATOM	1046	CA	ASP	132	-62.684	104.903	-19.526	1.00	34.31
60	ATOM	1047	CB	ASP	132	-62.420	103.837	-18.464	1.00	42.68
	ATOM	1048	CG	ASP	132	-63.405	103.903	-17.309	1.00	47.98
	ATOM	1049	OD1	ASP	132	-63.422	102.954	-16.485	1.00	50.41
	ATOM	1050	OD2	ASP	132	-64.152	104.909	-17.223	1.00	46.42
	ATOM	1051	C	ASP	132	-61.471	105.052	-20.428	1.00	34.32
65	ATOM	1052	O	ASP	132	-61.211	104.207	-21.278	1.00	32.87
	ATOM	1053	N	GLY	133	-60.735	106.138	-20.233	1.00	34.53

	ATOM	1054	CA	GLY	133	-59.570	106.407	-21.051	1.00	36.93
	ATOM	1055	C	GLY	133	-58.515	105.322	-21.139	1.00	35.06
	ATOM	1056	O	GLY	133	-57.735	105.292	-22.091	1.00	40.44
	ATOM	1057	N	ASP	134	-58.480	104.423	-20.166	1.00	31.08
5	ATOM	1058	CA	ASP	134	-57.475	103.379	-20.187	1.00	27.63
	ATOM	1059	CB	ASP	134	-56.978	103.102	-18.766	1.00	28.43
	ATOM	1060	CG	ASP	134	-58.076	102.644	-17.831	1.00	31.78
	ATOM	1061	OD1	ASP	134	-59.224	103.100	-17.986	1.00	35.68
	ATOM	1062	OD2	ASP	134	-57.786	101.837	-16.924	1.00	32.53
10	ATOM	1063	C	ASP	134	-57.917	102.100	-20.873	1.00	25.08
	ATOM	1064	O	ASP	134	-57.093	101.328	-21.310	1.00	19.71
	ATOM	1065	N	VAL	135	-59.215	101.882	-21.010	1.00	26.84
	ATOM	1066	CA	VAL	135	-59.666	100.661	-21.661	1.00	23.72
	ATOM	1067	CB	VAL	135	-60.820	100.013	-20.893	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-60.295	99.452	-19.600	1.00	22.76
	ATOM	1069	CG2	VAL	135	-61.916	101.024	-20.635	1.00	22.12
	ATOM	1070	C	VAL	135	-60.048	100.804	-23.131	1.00	26.54
	ATOM	1071	O	VAL	135	-59.994	99.829	-23.869	1.00	29.26
	ATOM	1072	N	THR	136	-60.442	101.998	-23.568	1.00	26.39
20	ATOM	1073	CA	THR	136	-60.763	102.175	-24.983	1.00	28.40
	ATOM	1074	CB	THR	136	-62.307	102.206	-25.239	1.00	28.21
	ATOM	1075	OG1	THR	136	-62.746	103.546	-25.455	1.00	37.13
	ATOM	1076	CG2	THR	136	-63.055	101.617	-24.069	1.00	26.90
	ATOM	1077	C	THR	136	-60.067	103.424	-25.544	1.00	31.60
25	ATOM	1078	O	THR	136	-60.315	104.551	-25.107	1.00	27.29
	ATOM	1079	N	PHE	137	-59.170	103.195	-26.505	1.00	31.45
	ATOM	1080	CA	PHE	137	-58.386	104.263	-27.113	1.00	26.64
	ATOM	1081	CB	PHE	137	-57.074	104.396	-26.352	1.00	26.84
	ATOM	1082	CG	PHE	137	-56.435	103.080	-26.014	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-55.598	102.441	-26.923	1.00	27.77
	ATOM	1084	CD2	PHE	137	-56.660	102.483	-24.785	1.00	23.70
	ATOM	1085	CE1	PHE	137	-54.996	101.234	-26.612	1.00	27.32
	ATOM	1086	CE2	PHE	137	-56.062	101.273	-24.467	1.00	26.31
	ATOM	1087	CZ	PHE	137	-55.227	100.650	-25.384	1.00	29.43
35	ATOM	1088	C	PHE	137	-58.141	104.036	-28.602	1.00	28.06
	ATOM	1089	O	PHE	137	-58.346	102.938	-29.100	1.00	25.04
	ATOM	1090	N	PHE	138	-57.696	105.078	-29.304	1.00	29.81
	ATOM	1091	CA	PHE	138	-57.488	105.006	-30.748	1.00	29.93
	ATOM	1092	CB	PHE	138	-58.610	105.797	-31.417	1.00	27.47
40	ATOM	1093	CG	PHE	138	-58.717	105.590	-32.890	1.00	28.49
	ATOM	1094	CD1	PHE	138	-58.222	104.437	-33.490	1.00	27.65
	ATOM	1095	CD2	PHE	138	-59.325	106.557	-33.686	1.00	31.06
	ATOM	1096	CE1	PHE	138	-58.330	104.255	-34.863	1.00	30.90
	ATOM	1097	CE2	PHE	138	-59.440	106.387	-35.060	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-58.941	105.235	-35.651	1.00	27.41
	ATOM	1099	C	PHE	138	-56.093	105.477	-31.206	1.00	30.91
	ATOM	1100	O	PHE	138	-55.644	106.568	-30.876	1.00	24.94
	ATOM	1101	N	GLY	139	-55.456	104.616	-32.004	1.00	36.94
	ATOM	1102	CA	GLY	139	-54.091	104.761	-32.507	1.00	36.08
50	ATOM	1103	C	GLY	139	-53.480	105.824	-33.386	1.00	36.48
	ATOM	1104	O	GLY	139	-53.632	107.002	-33.119	1.00	43.61
	ATOM	1105	N	ALA	140	-52.722	105.380	-34.390	1.00	36.60
	ATOM	1106	CA	ALA	140	-52.019	106.234	-35.368	1.00	34.94
	ATOM	1107	CB	ALA	140	-52.917	107.375	-35.811	1.00	36.94
55	ATOM	1108	C	ALA	140	-50.635	106.792	-34.985	1.00	32.53
	ATOM	1109	O	ALA	140	-50.514	107.733	-34.208	1.00	27.96
	ATOM	1110	N	LEU	141	-49.596	106.206	-35.576	1.00	33.23
	ATOM	1111	CA	LEU	141	-48.198	106.603	-35.358	1.00	36.11
	ATOM	1112	CB	LEU	141	-47.536	105.643	-34.371	1.00	37.53
60	ATOM	1113	CG	LEU	141	-46.048	105.764	-34.025	1.00	38.87
	ATOM	1114	CD1	LEU	141	-45.751	104.868	-32.848	1.00	40.67
	ATOM	1115	CD2	LEU	141	-45.172	105.365	-35.196	1.00	37.53
	ATOM	1116	C	LEU	141	-47.467	106.547	-36.704	1.00	38.89
	ATOM	1117	O	LEU	141	-47.531	105.538	-37.403	1.00	39.25
65	ATOM	1118	N	LYS	142	-46.764	107.612	-37.074	1.00	42.86
	ATOM	1119	CA	LYS	142	-46.070	107.601	-38.357	1.00	42.96

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	ATOM	1120	CB	LYS	142	-45.935	109.021	-38.899	1.00	41.80
	ATOM	1121	CG	LYS	142	-45.196	109.061	-40.216	1.00	46.88
	ATOM	1122	CD	LYS	142	-45.453	110.335	-40.992	1.00	49.58
	ATOM	1123	CE	LYS	142	-44.680	110.297	-42.300	1.00	51.07
5	ATOM	1124	NZ	LYS	142	-44.950	111.480	-43.154	1.00	56.20
	ATOM	1125	C	LYS	142	-44.703	106.927	-38.346	1.00	43.41
	ATOM	1126	O	LYS	142	-43.856	107.231	-37.511	1.00	44.98
	ATOM	1127	N	LEU	143	-44.492	106.012	-39.287	1.00	42.81
	ATOM	1128	CA	LEU	143	-43.225	105.293	-39.397	1.00	40.65
10	ATOM	1129	CB	LEU	143	-43.436	103.948	-40.088	1.00	36.56
	ATOM	1130	CG	LEU	143	-44.436	102.966	-39.479	1.00	36.17
	ATOM	1131	CD1	LEU	143	-44.525	101.734	-40.361	1.00	37.95
	ATOM	1132	CD2	LEU	143	-44.005	102.587	-38.079	1.00	27.38
	ATOM	1133	C	LEU	143	-42.222	106.102	-40.211	1.00	43.14
15	ATOM	1134	O	LEU	143	-42.604	106.850	-41.117	1.00	46.28
	ATOM	1135	N	LEU	144	-40.942	105.957	-39.895	1.00	42.63
	ATOM	1136	CA	LEU	144	-39.913	106.676	-40.632	1.00	44.11
	ATOM	1137	CB	LEU	144	-38.623	106.749	-39.821	1.00	44.88
	ATOM	1138	CG	LEU	144	-38.731	107.564	-38.534	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-37.440	107.487	-37.766	1.00	47.71
	ATOM	1140	CD2	LEU	144	-39.062	109.008	-38.868	1.00	47.87
	ATOM	1141	C	LEU	144	-39.637	105.975	-41.947	1.00	45.66
	ATOM	1142	O	LEU	144	-39.961	104.778	-42.058	1.00	46.37
	ATOM	1143	OXT	LEU	144	-39.085	106.633	-42.849	1.00	50.66
25	END					-118.331	85.231	-119.150	0.00	0.00

TABLE 12

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	ATOM	1	CB	VAL	1	-4.730	130.789	-9.142	1.00 69.28
5	ATOM	2	CG1	VAL	1	-5.405	132.112	-9.568	1.00 68.26
	ATOM	3	CG2	VAL	1	-4.224	130.008	-10.360	1.00 71.75
	ATOM	4	C	VAL	1	-4.154	131.490	-6.801	1.00 64.86
	ATOM	5	O	VAL	1	-4.144	132.677	-6.440	1.00 64.02
	ATOM	6	N	VAL	1	-2.665	129.884	-8.004	1.00 65.10
10	ATOM	7	CA	VAL	1	-3.551	131.078	-8.158	1.00 65.98
	ATOM	8	N	THR	2	-4.685	130.515	-6.059	1.00 61.42
	ATOM	9	CA	THR	2	-5.280	130.791	-4.752	1.00 56.64
	ATOM	10	CB	THR	2	-6.767	130.396	-4.702	1.00 56.77
	ATOM	11	OG1	THR	2	-6.897	129.021	-5.073	1.00 57.45
15	ATOM	12	CG2	THR	2	-7.596	131.260	-5.644	1.00 55.68
	ATOM	13	C	THR	2	-4.581	130.026	-3.646	1.00 53.08
	ATOM	14	O	THR	2	-3.775	129.137	-3.900	1.00 52.00
	ATOM	15	N	GLN	3	-4.913	130.370	-2.408	1.00 50.34
	ATOM	16	CA	GLN	3	-4.319	129.715	-1.249	1.00 45.97
20	ATOM	17	CB	GLN	3	-3.743	130.759	-0.298	1.00 46.09
	ATOM	18	CG	GLN	3	-2.872	131.785	-0.973	1.00 49.60
	ATOM	19	CD	GLN	3	-2.261	132.745	0.015	1.00 50.43
	ATOM	20	OE1	GLN	3	-1.422	132.356	0.821	1.00 50.94
	ATOM	21	NE2	GLN	3	-2.684	134.006	-0.034	1.00 49.99
25	ATOM	22	C	GLN	3	-5.350	128.883	-0.509	1.00 42.41
	ATOM	23	O	GLN	3	-6.259	129.412	0.124	1.00 37.34
	ATOM	24	N	ASP	4	-5.205	127.571	-0.594	1.00 40.32
	ATOM	25	CA	ASP	4	-6.130	126.675	0.076	1.00 39.37
	ATOM	26	CB	ASP	4	-5.874	125.230	-0.359	1.00 44.56
30	ATOM	27	CG	ASP	4	-6.176	125.001	-1.825	1.00 45.76
	ATOM	28	OD1	ASP	4	-6.227	126.001	-2.578	1.00 48.58
	ATOM	29	OD2	ASP	4	-6.352	123.830	-2.223	1.00 45.55
	ATOM	30	C	ASP	4	-5.940	126.795	1.572	1.00 36.30
	ATOM	31	O	ASP	4	-4.834	127.034	2.050	1.00 36.29
35	ATOM	32	N	CYS	5	-7.027	126.642	2.310	1.00 35.11
	ATOM	33	CA	CYS	5	-6.971	126.698	3.759	1.00 35.61
	ATOM	34	CB	CYS	5	-6.848	128.142	4.252	1.00 34.96
	ATOM	35	SG	CYS	5	-7.880	129.325	3.412	1.00 35.34
	ATOM	36	C	CYS	5	-8.200	126.032	4.342	1.00 31.80
40	ATOM	37	O	CYS	5	-9.252	125.993	3.717	1.00 31.85
	ATOM	38	N	LEU	6	-8.042	125.472	5.531	1.00 27.72
	ATOM	39	CA	LEU	6	-9.129	124.808	6.219	1.00 30.80
	ATOM	40	CB	LEU	6	-9.008	123.296	6.062	1.00 29.25
	ATOM	41	CG	LEU	6	-10.031	122.421	6.788	1.00 29.47
45	ATOM	42	CD1	LEU	6	-10.164	121.101	6.078	1.00 30.65
	ATOM	43	CD2	LEU	6	-9.593	122.196	8.204	1.00 35.27
	ATOM	44	C	LEU	6	-9.037	125.197	7.681	1.00 31.83
	ATOM	45	O	LEU	6	-7.981	125.069	8.289	1.00 33.45
	ATOM	46	N	GLN	7	-10.136	125.685	8.244	1.00 32.23
50	ATOM	47	CA	GLN	7	-10.140	126.099	9.631	1.00 30.26
	ATOM	48	CB	GLN	7	-10.467	127.585	9.721	1.00 29.56
	ATOM	49	CG	GLN	7	-10.218	128.212	11.075	1.00 25.09
	ATOM	50	CD	GLN	7	-10.317	129.724	11.018	1.00 26.50
	ATOM	51	OE1	GLN	7	-11.400	130.288	10.863	1.00 24.40
55	ATOM	52	NE2	GLN	7	-9.180	130.387	11.127	1.00 27.03
	ATOM	53	C	GLN	7	-11.146	125.288	10.418	1.00 32.11
	ATOM	54	O	GLN	7	-12.229	124.982	9.935	1.00 34.24
	ATOM	55	N	LEU	8	-10.761	124.929	11.634	1.00 33.73
	ATOM	56	CA	LEU	8	-11.604	124.148	12.524	1.00 35.41
60	ATOM	57	CB	LEU	8	-10.879	122.855	12.935	1.00 32.44
	ATOM	58	CG	LEU	8	-11.088	121.554	12.145	1.00 27.69
	ATOM	59	CD1	LEU	8	-11.652	121.817	10.774	1.00 26.43
	ATOM	60	CD2	LEU	8	-9.772	120.828	12.071	1.00 18.64
	ATOM	61	C	LEU	8	-11.962	124.966	13.756	1.00 36.87
65	ATOM	62	O	LEU	8	-11.181	125.792	14.226	1.00 36.05
	ATOM	63	N	ILE	9	-13.154	124.716	14.277	1.00 39.25

	ATOM	64	CA	ILE	9	-13.668	125.420	15.440	1.00	35.57
	ATOM	65	CB	ILE	9	-14.863	126.302	14.998	1.00	34.28
	ATOM	66	CG2	ILE	9	-16.034	126.169	15.938	1.00	38.20
	ATOM	67	CG1	ILE	9	-14.417	127.741	14.893	1.00	35.22
5	ATOM	68	CD1	ILE	9	-15.549	128.640	14.527	1.00	43.63
	ATOM	69	C	ILE	9	-14.088	124.406	16.508	1.00	35.28
	ATOM	70	O	ILE	9	-14.535	123.309	16.184	1.00	33.75
	ATOM	71	N	ALA	10	-13.932	124.759	17.778	1.00	35.78
	ATOM	72	CA	ALA	10	-14.322	123.849	18.844	1.00	35.28
10	ATOM	73	CB	ALA	10	-13.944	124.424	20.189	1.00	28.47
	ATOM	74	C	ALA	10	-15.821	123.583	18.797	1.00	36.49
	ATOM	75	O	ALA	10	-16.618	124.486	18.561	1.00	37.41
	ATOM	76	N	ASP	11	-16.198	122.332	19.020	1.00	39.44
	ATOM	77	CA	ASP	11	-17.603	121.943	19.013	1.00	40.77
15	ATOM	78	CB	ASP	11	-17.788	120.667	18.191	1.00	38.90
	ATOM	79	CG	ASP	11	-19.201	120.141	18.254	1.00	40.34
	ATOM	80	OD1	ASP	11	-20.134	120.946	18.490	1.00	39.50
	ATOM	81	OD2	ASP	11	-19.376	118.924	18.056	1.00	37.38
	ATOM	82	C	ASP	11	-18.094	121.728	20.440	1.00	42.25
20	ATOM	83	O	ASP	11	-17.956	120.643	21.007	1.00	40.54
	ATOM	84	N	SER	12	-18.675	122.777	21.008	1.00	42.92
	ATOM	85	CA	SER	12	-19.167	122.757	22.384	1.00	43.84
	ATOM	86	CB	SER	12	-19.646	124.150	22.776	1.00	42.13
	ATOM	87	OG	SER	12	-20.662	124.590	21.893	1.00	43.31
25	ATOM	88	C	SER	12	-20.281	121.761	22.659	1.00	46.10
	ATOM	89	O	SER	12	-20.704	121.599	23.804	1.00	43.90
	ATOM	90	N	GLU	13	-20.750	121.087	21.616	1.00	46.72
	ATOM	91	CA	GLU	13	-21.825	120.128	21.782	1.00	44.93
	ATOM	92	CB	GLU	13	-22.870	120.333	20.701	1.00	46.71
30	ATOM	93	CG	GLU	13	-23.580	121.649	20.856	1.00	58.25
	ATOM	94	CD	GLU	13	-24.956	121.636	20.224	1.00	66.70
	ATOM	95	OE1	GLU	13	-25.778	120.769	20.633	1.00	74.10
	ATOM	96	OE2	GLU	13	-25.221	122.481	19.327	1.00	71.55
	ATOM	97	C	GLU	13	-21.412	118.661	21.855	1.00	43.24
35	ATOM	98	O	GLU	13	-22.256	117.768	21.751	1.00	44.25
	ATOM	99	N	THR	14	-20.122	118.406	22.019	1.00	37.06
	ATOM	100	CA	THR	14	-19.660	117.032	22.175	1.00	36.77
	ATOM	101	CB	THR	14	-19.145	116.407	20.844	1.00	33.39
	ATOM	102	OG1	THR	14	-17.866	116.946	20.513	1.00	41.69
40	ATOM	103	CG2	THR	14	-20.118	116.686	19.710	1.00	31.66
	ATOM	104	C	THR	14	-18.544	117.070	23.213	1.00	34.96
	ATOM	105	O	THR	14	-17.847	118.068	23.351	1.00	35.04
	ATOM	106	N	PRO	15	-18.379	115.987	23.973	1.00	37.11
	ATOM	107	CD	PRO	15	-19.178	114.754	23.910	1.00	36.50
45	ATOM	108	CA	PRO	15	-17.352	115.892	25.015	1.00	38.81
	ATOM	109	CB	PRO	15	-17.646	114.542	25.669	1.00	39.15
	ATOM	110	CG	PRO	15	-19.090	114.276	25.322	1.00	37.32
	ATOM	111	C	PRO	15	-15.935	115.938	24.459	1.00	37.78
	ATOM	112	O	PRO	15	-15.703	115.533	23.323	1.00	35.91
50	ATOM	113	N	THR	16	-14.984	116.423	25.251	1.00	36.67
	ATOM	114	CA	THR	16	-13.609	116.463	24.783	1.00	38.46
	ATOM	115	CB	THR	16	-12.684	117.217	25.757	1.00	38.61
	ATOM	116	OG1	THR	16	-12.634	116.523	27.008	1.00	42.44
	ATOM	117	CG2	THR	16	-13.196	118.631	25.985	1.00	40.11
55	ATOM	118	C	THR	16	-13.158	115.020	24.708	1.00	35.97
	ATOM	119	O	THR	16	-13.465	114.237	25.597	1.00	39.85
	ATOM	120	N	ILE	17	-12.439	114.665	23.651	1.00	33.66
	ATOM	121	CA	ILE	17	-11.970	113.298	23.484	1.00	34.76
	ATOM	122	CB	ILE	17	-11.565	113.038	22.028	1.00	32.44
60	ATOM	123	CG2	ILE	17	-10.965	111.652	21.892	1.00	28.66
	ATOM	124	CG1	ILE	17	-12.788	113.209	21.120	1.00	33.15
	ATOM	125	CD1	ILE	17	-12.498	113.055	19.640	1.00	31.19
	ATOM	126	C	ILE	17	-10.797	112.944	24.389	1.00	40.64
	ATOM	127	O	ILE	17	-9.793	113.644	24.419	1.00	44.07
65	ATOM	128	N	GLN	18	-10.929	111.848	25.131	1.00	45.91
	ATOM	129	CA	GLN	18	-9.868	111.396	26.028	1.00	50.22

	ATOM	130	CB	GLN	18	-10.437	111.010	27.389	1.00	53.09
	ATOM	131	CG	GLN	18	-9.809	111.778	28.522	1.00	57.60
	ATOM	132	CD	GLN	18	-10.197	113.233	28.470	1.00	61.39
	ATOM	133	OE1	GLN	18	-11.356	113.586	28.705	1.00	68.36
5	ATOM	134	NE2	GLN	18	-9.239	114.089	28.141	1.00	64.93
	ATOM	135	C	GLN	18	-9.151	110.192	25.445	1.00	51.56
	ATOM	136	O	GLN	18	-9.783	109.241	24.995	1.00	56.44
	ATOM	137	N	LYS	19	-7.829	110.227	25.459	1.00	52.60
	ATOM	138	CA	LYS	19	-7.066	109.113	24.922	1.00	56.10
10	ATOM	139	CB	LYS	19	-7.217	109.042	23.401	1.00	55.55
	ATOM	140	CG	LYS	19	-6.348	107.961	22.800	1.00	61.15
	ATOM	141	CD	LYS	19	-6.633	107.715	21.333	1.00	64.42
	ATOM	142	CE	LYS	19	-5.728	106.595	20.800	1.00	64.84
	ATOM	143	NZ	LYS	19	-5.986	106.276	19.359	1.00	69.52
15	ATOM	144	C	LYS	19	-5.589	109.187	25.278	1.00	57.41
	ATOM	145	O	LYS	19	-4.970	110.247	25.164	1.00	61.02
	ATOM	146	N	GLY	20	-5.034	108.053	25.703	1.00	58.84
	ATOM	147	CA	GLY	20	-3.632	107.989	26.072	1.00	56.66
	ATOM	148	C	GLY	20	-3.214	109.103	27.011	1.00	56.87
20	ATOM	149	O	GLY	20	-2.105	109.626	26.889	1.00	58.64
	ATOM	150	N	SER	21	-4.097	109.462	27.945	1.00	54.49
	ATOM	151	CA	SER	21	-3.835	110.524	28.919	1.00	53.68
	ATOM	152	CB	SER	21	-2.623	110.166	29.790	1.00	55.80
	ATOM	153	OG	SER	21	-1.404	110.534	29.172	1.00	65.34
25	ATOM	154	C	SER	21	-3.629	111.906	28.252	1.00	51.41
	ATOM	155	O	SER	21	-2.944	112.793	28.790	1.00	47.74
	ATOM	156	N	TYR	22	-4.229	112.066	27.074	1.00	46.25
	ATOM	157	CA	TYR	22	-4.176	113.308	26.315	1.00	40.57
	ATOM	158	CB	TYR	22	-3.507	113.083	24.957	1.00	43.27
30	ATOM	159	CG	TYR	22	-2.007	113.220	24.975	1.00	49.18
	ATOM	160	CD1	TYR	22	-1.222	112.408	25.787	1.00	52.61
	ATOM	161	CE1	TYR	22	0.169	112.542	25.817	1.00	51.92
	ATOM	162	CD2	TYR	22	-1.367	114.172	24.189	1.00	50.01
	ATOM	163	CE2	TYR	22	0.021	114.311	24.209	1.00	50.94
35	ATOM	164	CZ	TYR	22	0.782	113.494	25.026	1.00	50.78
	ATOM	165	OH	TYR	22	2.152	113.634	25.049	1.00	53.87
	ATOM	166	C	TYR	22	-5.610	113.777	26.095	1.00	37.85
	ATOM	167	O	TYR	22	-6.525	112.959	26.003	1.00	38.70
	ATOM	168	N	THR	23	-5.816	115.085	26.029	1.00	33.81
40	ATOM	169	CA	THR	23	-7.155	115.612	25.787	1.00	32.40
	ATOM	170	CB	THR	23	-7.535	116.753	26.765	1.00	31.58
	ATOM	171	OG1	THR	23	-7.427	116.299	28.118	1.00	30.76
	ATOM	172	CG2	THR	23	-8.956	117.207	26.507	1.00	24.08
	ATOM	173	C	THR	23	-7.181	116.177	24.378	1.00	33.28
45	ATOM	174	O	THR	23	-6.288	116.923	23.991	1.00	30.44
	ATOM	175	N	PHE	24	-8.202	115.818	23.615	1.00	32.16
	ATOM	176	CA	PHE	24	-8.334	116.307	22.255	1.00	30.68
	ATOM	177	CB	PHE	24	-8.317	115.145	21.262	1.00	28.20
	ATOM	178	CG	PHE	24	-7.007	114.425	21.204	1.00	31.00
50	ATOM	179	CD1	PHE	24	-6.709	113.425	22.115	1.00	31.69
	ATOM	180	CD2	PHE	24	-6.052	114.769	20.252	1.00	31.42
	ATOM	181	CE1	PHE	24	-5.479	112.780	22.080	1.00	32.57
	ATOM	182	CE2	PHE	24	-4.825	114.130	20.211	1.00	25.01
	ATOM	183	CZ	PHE	24	-4.538	113.134	21.128	1.00	30.12
55	ATOM	184	C	PHE	24	-9.609	117.121	22.069	1.00	31.73
	ATOM	185	O	PHE	24	-10.697	116.691	22.440	1.00	33.31
	ATOM	186	N	VAL	25	-9.463	118.305	21.498	1.00	32.92
	ATOM	187	CA	VAL	25	-10.596	119.170	21.253	1.00	30.84
	ATOM	188	CB	VAL	25	-10.137	120.562	20.791	1.00	30.29
60	ATOM	189	CG1	VAL	25	-11.333	121.433	20.501	1.00	28.13
	ATOM	190	CG2	VAL	25	-9.264	121.191	21.842	1.00	29.53
	ATOM	191	C	VAL	25	-11.478	118.577	20.159	1.00	36.08
	ATOM	192	O	VAL	25	-10.981	118.083	19.149	1.00	35.82
	ATOM	193	N	PRO	26	-12.803	118.595	20.360	1.00	38.03
65	ATOM	194	CD	PRO	26	-13.477	118.871	21.638	1.00	38.28
	ATOM	195	CA	PRO	26	-13.761	118.065	19.379	1.00	38.08

	ATOM	196	CB	PRO	26	-15.043	117.888	20.196	1.00	37.62
	ATOM	197	CG	PRO	26	-14.579	117.861	21.617	1.00	36.31
	ATOM	198	C	PRO	26	-13.930	119.158	18.324	1.00	40.85
	ATOM	199	O	PRO	26	-14.487	120.215	18.612	1.00	42.27
5	ATOM	200	N	TRP	27	-13.459	118.922	17.108	1.00	40.86
	ATOM	201	CA	TRP	27	-13.561	119.960	16.090	1.00	37.05
	ATOM	202	CB	TRP	27	-12.361	119.897	15.142	1.00	33.29
	ATOM	203	CG	TRP	27	-11.056	120.095	15.824	1.00	31.06
	ATOM	204	CD2	TRP	27	-10.633	121.253	16.546	1.00	27.79
10	ATOM	205	CE2	TRP	27	-9.349	120.984	17.046	1.00	29.28
	ATOM	206	CE3	TRP	27	-11.216	122.491	16.820	1.00	27.06
	ATOM	207	CD1	TRP	27	-10.041	119.201	15.908	1.00	31.07
	ATOM	208	NE1	TRP	27	-9.012	119.721	16.640	1.00	32.70
	ATOM	209	CZ2	TRP	27	-8.630	121.907	17.810	1.00	25.99
15	ATOM	210	CZ3	TRP	27	-10.502	123.411	17.580	1.00	26.93
	ATOM	211	CH2	TRP	27	-9.222	123.112	18.065	1.00	28.08
	ATOM	212	C	TRP	27	-14.832	119.957	15.272	1.00	38.50
	ATOM	213	O	TRP	27	-15.523	118.949	15.165	1.00	39.07
	ATOM	214	N	LEU	28	-15.121	121.121	14.702	1.00	40.19
20	ATOM	215	CA	LEU	28	-16.278	121.335	13.848	1.00	39.45
	ATOM	216	CB	LEU	28	-17.351	122.116	14.596	1.00	44.18
	ATOM	217	CG	LEU	28	-18.748	122.083	13.980	1.00	46.83
	ATOM	218	CD1	LEU	28	-19.301	120.658	14.088	1.00	45.15
	ATOM	219	CD2	LEU	28	-19.660	123.076	14.707	1.00	49.15
25	ATOM	220	C	LEU	28	-15.737	122.185	12.706	1.00	39.80
	ATOM	221	O	LEU	28	-14.948	123.089	12.940	1.00	42.40
	ATOM	222	N	LEU	29	-16.140	121.903	11.479	1.00	33.58
	ATOM	223	CA	LEU	29	-15.637	122.687	10.367	1.00	29.21
	ATOM	224	CB	LEU	29	-16.098	122.101	9.034	1.00	27.16
30	ATOM	225	CG	LEU	29	-15.664	122.913	7.810	1.00	22.36
	ATOM	226	CD1	LEU	29	-14.177	122.711	7.566	1.00	21.47
	ATOM	227	CD2	LEU	29	-16.460	122.504	6.600	1.00	14.53
	ATOM	228	C	LEU	29	-16.074	124.145	10.436	1.00	29.98
	ATOM	229	O	LEU	29	-17.255	124.453	10.560	1.00	30.36
35	ATOM	230	N	SER	30	-15.098	125.040	10.358	1.00	29.88
	ATOM	231	CA	SER	30	-15.369	126.465	10.364	1.00	28.49
	ATOM	232	CB	SER	30	-14.192	127.241	10.955	1.00	26.13
	ATOM	233	OG	SER	30	-14.418	128.630	10.873	1.00	20.25
	ATOM	234	C	SER	30	-15.555	126.818	8.894	1.00	31.61
40	ATOM	235	O	SER	30	-16.560	127.406	8.507	1.00	35.14
	ATOM	236	N	PHE	31	-14.577	126.442	8.077	1.00	32.12
	ATOM	237	CA	PHE	31	-14.644	126.685	6.646	1.00	30.76
	ATOM	238	CB	PHE	31	-14.559	128.186	6.341	1.00	27.55
	ATOM	239	CG	PHE	31	-13.158	128.708	6.186	1.00	31.23
45	ATOM	240	CD1	PHE	31	-12.482	128.577	4.978	1.00	32.00
	ATOM	241	CD2	PHE	31	-12.506	129.321	7.254	1.00	27.89
	ATOM	242	CE1	PHE	31	-11.185	129.045	4.836	1.00	29.70
	ATOM	243	CE2	PHE	31	-11.210	129.791	7.116	1.00	26.15
	ATOM	244	CZ	PHE	31	-10.548	129.653	5.908	1.00	28.30
50	ATOM	245	C	PHE	31	-13.522	125.940	5.943	1.00	32.06
	ATOM	246	O	PHE	31	-12.459	125.697	6.514	1.00	32.26
	ATOM	247	N	LYS	32	-13.776	125.563	4.702	1.00	34.83
	ATOM	248	CA	LYS	32	-12.791	124.861	3.906	1.00	37.13
	ATOM	249	CB	LYS	32	-13.219	123.410	3.699	1.00	36.55
55	ATOM	250	CG	LYS	32	-12.278	122.629	2.824	1.00	40.62
	ATOM	251	CD	LYS	32	-12.862	121.301	2.409	1.00	44.56
	ATOM	252	CE	LYS	32	-12.005	120.702	1.292	1.00	46.29
	ATOM	253	NZ	LYS	32	-12.432	119.326	0.899	1.00	50.58
	ATOM	254	C	LYS	32	-12.703	125.590	2.568	1.00	36.39
60	ATOM	255	O	LYS	32	-13.726	125.923	1.968	1.00	38.69
	ATOM	256	N	ARG	33	-11.484	125.857	2.112	1.00	37.90
	ATOM	257	CA	ARG	33	-11.290	126.557	0.850	1.00	35.74
	ATOM	258	CB	ARG	33	-10.861	127.992	1.134	1.00	36.79
	ATOM	259	CG	ARG	33	-10.542	128.830	-0.083	1.00	31.09
65	ATOM	260	CD	ARG	33	-10.402	130.291	0.325	1.00	31.62
	ATOM	261	NE	ARG	33	-10.243	131.198	-0.809	1.00	38.16

	ATOM	262	CZ	ARG	33	-9.080	131.703	-1.204	1.00	39.15
	ATOM	263	NH1	ARG	33	-7.966	131.394	-0.555	1.00	43.41
	ATOM	264	NH2	ARG	33	-9.026	132.513	-2.248	1.00	40.65
	ATOM	265	C	ARG	33	-10.238	125.842	0.011	1.00	37.71
5	ATOM	266	O	ARG	33	-9.084	125.701	0.429	1.00	41.16
	ATOM	267	N	GLY	34	-10.628	125.378	-1.170	1.00	36.59
	ATOM	268	CA	GLY	34	-9.678	124.688	-2.018	1.00	36.72
	ATOM	269	C	GLY	34	-9.692	123.188	-1.812	1.00	40.01
	ATOM	270	O	GLY	34	-10.603	122.638	-1.194	1.00	39.85
10	ATOM	271	N	SER	35	-8.657	122.529	-2.313	1.00	38.74
	ATOM	272	CA	SER	35	-8.556	121.079	-2.226	1.00	40.23
	ATOM	273	CB	SER	35	-8.506	120.511	-3.639	1.00	43.02
	ATOM	274	OG	SER	35	-7.463	121.135	-4.386	1.00	48.05
	ATOM	275	C	SER	35	-7.361	120.536	-1.439	1.00	40.54
15	ATOM	276	O	SER	35	-7.367	119.369	-1.049	1.00	43.74
	ATOM	277	N	ALA	36	-6.349	121.368	-1.208	1.00	35.56
	ATOM	278	CA	ALA	36	-5.142	120.942	-0.507	1.00	29.62
	ATOM	279	CB	ALA	36	-4.142	122.080	-0.491	1.00	25.47
	ATOM	280	C	ALA	36	-5.331	120.402	0.908	1.00	31.23
20	ATOM	281	O	ALA	36	-4.512	119.625	1.381	1.00	29.56
	ATOM	282	N	LEU	37	-6.401	120.805	1.585	1.00	34.09
	ATOM	283	CA	LEU	37	-6.640	120.362	2.955	1.00	32.32
	ATOM	284	CB	LEU	37	-6.378	121.513	3.922	1.00	26.47
	ATOM	285	CG	LEU	37	-4.960	122.078	3.881	1.00	25.72
25	ATOM	286	CD1	LEU	37	-4.925	123.457	4.491	1.00	24.61
	ATOM	287	CD2	LEU	37	-4.019	121.140	4.600	1.00	27.53
	ATOM	288	C	LEU	37	-8.044	119.827	3.173	1.00	35.16
	ATOM	289	O	LEU	37	-9.012	120.344	2.626	1.00	36.47
	ATOM	290	N	GLU	38	-8.143	118.789	3.988	1.00	38.30
30	ATOM	291	CA	GLU	38	-9.422	118.170	4.297	1.00	41.17
	ATOM	292	CB	GLU	38	-9.587	116.872	3.516	1.00	46.15
	ATOM	293	CG	GLU	38	-10.207	117.008	2.146	1.00	50.01
	ATOM	294	CD	GLU	38	-10.074	115.723	1.340	1.00	51.18
	ATOM	295	OE1	GLU	38	-10.234	114.630	1.941	1.00	48.40
35	ATOM	296	OE2	GLU	38	-9.809	115.810	0.111	1.00	50.23
	ATOM	297	C	GLU	38	-9.524	117.836	5.773	1.00	42.69
	ATOM	298	O	GLU	38	-8.522	117.687	6.460	1.00	42.55
	ATOM	299	N	GLU	39	-10.746	117.716	6.265	1.00	44.75
	ATOM	300	CA	GLU	39	-10.937	117.356	7.660	1.00	46.50
40	ATOM	301	CB	GLU	39	-12.173	118.045	8.228	1.00	47.52
	ATOM	302	CG	GLU	39	-13.318	118.109	7.241	1.00	58.50
	ATOM	303	CD	GLU	39	-14.587	118.669	7.856	1.00	62.35
	ATOM	304	OE1	GLU	39	-15.546	118.955	7.094	1.00	64.58
	ATOM	305	OE2	GLU	39	-14.625	118.811	9.101	1.00	63.04
45	ATOM	306	C	GLU	39	-11.135	115.852	7.628	1.00	44.56
	ATOM	307	O	GLU	39	-11.824	115.336	6.757	1.00	48.11
	ATOM	308	N	LYS	40	-10.515	115.144	8.558	1.00	40.71
	ATOM	309	CA	LYS	40	-10.636	113.700	8.595	1.00	38.08
	ATOM	310	CB	LYS	40	-9.576	113.060	7.702	1.00	41.28
50	ATOM	311	CG	LYS	40	-9.506	111.542	7.812	1.00	41.52
	ATOM	312	CD	LYS	40	-8.234	111.007	7.167	1.00	42.73
	ATOM	313	CE	LYS	40	-8.056	109.517	7.400	1.00	41.97
	ATOM	314	NZ	LYS	40	-6.741	109.048	6.864	1.00	45.94
	ATOM	315	C	LYS	40	-10.483	113.175	10.006	1.00	39.66
55	ATOM	316	O	LYS	40	-9.414	113.267	10.604	1.00	36.18
	ATOM	317	N	GLU	41	-11.568	112.623	10.534	1.00	40.16
	ATOM	318	CA	GLU	41	-11.575	112.051	11.872	1.00	42.80
	ATOM	319	CB	GLU	41	-10.857	110.707	11.836	1.00	45.31
	ATOM	320	CG	GLU	41	-11.457	109.772	10.792	1.00	56.07
60	ATOM	321	CD	GLU	41	-10.539	108.612	10.418	1.00	59.42
	ATOM	322	OE1	GLU	41	-9.373	108.861	10.021	1.00	63.71
	ATOM	323	OE2	GLU	41	-10.988	107.448	10.514	1.00	63.71
	ATOM	324	C	GLU	41	-10.944	112.974	12.902	1.00	38.93
	ATOM	325	O	GLU	41	-10.031	112.585	13.624	1.00	39.03
65	ATOM	326	N	ASN	42	-11.435	114.205	12.942	1.00	34.69
	ATOM	327	CA	ASN	42	-10.971	115.206	13.888	1.00	32.96

	ATOM	328	CB	ASN	42	-11.093	114.675	15.311	1.00	32.65
	ATOM	329	CG	ASN	42	-11.450	115.751	16.288	1.00	35.50
	ATOM	330	OD1	ASN	42	-10.886	115.829	17.371	1.00	40.26
	ATOM	331	ND2	ASN	42	-12.402	116.592	15.913	1.00	29.28
5	ATOM	332	C	ASN	42	-9.550	115.702	13.669	1.00	31.44
	ATOM	333	O	ASN	42	-8.949	116.288	14.564	1.00	22.78
	ATOM	334	N	LYS	43	-9.020	115.472	12.478	1.00	33.26
	ATOM	335	CA	LYS	43	-7.670	115.901	12.153	1.00	33.68
	ATOM	336	CB	LYS	43	-6.725	114.701	12.131	1.00	33.55
10	ATOM	337	CG	LYS	43	-6.507	114.071	13.484	1.00	40.49
	ATOM	338	CD	LYS	43	-5.644	112.840	13.374	1.00	45.39
	ATOM	339	CE	LYS	43	-6.352	111.756	12.586	1.00	52.57
	ATOM	340	NZ	LYS	43	-5.557	110.501	12.561	1.00	62.06
	ATOM	341	C	LYS	43	-7.663	116.569	10.797	1.00	32.05
15	ATOM	342	O	LYS	43	-8.625	116.462	10.048	1.00	37.32
	ATOM	343	N	ILE	44	-6.587	117.272	10.481	1.00	31.31
	ATOM	344	CA	ILE	44	-6.503	117.916	9.186	1.00	29.52
	ATOM	345	CB	ILE	44	-5.917	119.324	9.296	1.00	26.38
	ATOM	346	CG2	ILE	44	-5.804	119.940	7.917	1.00	26.09
20	ATOM	347	CG1	ILE	44	-6.829	120.188	10.168	1.00	28.61
	ATOM	348	CD1	ILE	44	-6.370	121.604	10.331	1.00	21.51
	ATOM	349	C	ILE	44	-5.640	117.061	8.275	1.00	31.60
	ATOM	350	O	ILE	44	-4.487	116.768	8.586	1.00	34.50
	ATOM	351	N	LEU	45	-6.216	116.644	7.156	1.00	31.44
25	ATOM	352	CA	LEU	45	-5.516	115.815	6.187	1.00	31.40
	ATOM	353	CB	LEU	45	-6.460	114.741	5.654	1.00	33.60
	ATOM	354	CG	LEU	45	-5.909	113.858	4.538	1.00	34.36
	ATOM	355	CD1	LEU	45	-4.780	112.995	5.063	1.00	29.00
	ATOM	356	CD2	LEU	45	-7.033	112.998	3.993	1.00	32.42
30	ATOM	357	C	LEU	45	-4.960	116.630	5.023	1.00	27.80
	ATOM	358	O	LEU	45	-5.670	117.414	4.401	1.00	27.94
	ATOM	359	N	VAL	46	-3.679	116.433	4.738	1.00	28.00
	ATOM	360	CA	VAL	46	-3.013	117.128	3.646	1.00	31.53
	ATOM	361	CB	VAL	46	-1.511	117.268	3.948	1.00	27.89
35	ATOM	362	CG1	VAL	46	-0.820	118.032	2.845	1.00	26.10
	ATOM	363	CG2	VAL	46	-1.327	117.964	5.268	1.00	23.39
	ATOM	364	C	VAL	46	-3.202	116.335	2.342	1.00	33.94
	ATOM	365	O	VAL	46	-2.863	115.152	2.276	1.00	39.68
	ATOM	366	N	LYS	47	-3.738	116.985	1.312	1.00	33.59
40	ATOM	367	CA	LYS	47	-3.968	116.318	0.033	1.00	37.05
	ATOM	368	CB	LYS	47	-5.402	116.571	-0.445	1.00	34.49
	ATOM	369	CG	LYS	47	-6.455	116.319	0.619	1.00	37.53
	ATOM	370	CD	LYS	47	-6.378	114.898	1.162	1.00	41.69
	ATOM	371	CE	LYS	47	-7.343	113.977	0.437	1.00	42.95
45	ATOM	372	NZ	LYS	47	-7.261	114.149	-1.031	1.00	44.50
	ATOM	373	C	LYS	47	-2.981	116.735	-1.058	1.00	37.34
	ATOM	374	O	LYS	47	-2.915	116.110	-2.111	1.00	43.04
	ATOM	375	N	GLU	48	-2.224	117.792	-0.801	1.00	36.99
	ATOM	376	CA	GLU	48	-1.225	118.288	-1.744	1.00	37.31
50	ATOM	377	CB	GLU	48	-1.706	119.551	-2.436	1.00	36.53
	ATOM	378	CG	GLU	48	-2.843	119.366	-3.397	1.00	46.74
	ATOM	379	CD	GLU	48	-3.281	120.689	-3.999	1.00	52.39
	ATOM	380	OE1	GLU	48	-2.386	121.520	-4.319	1.00	51.33
	ATOM	381	OE2	GLU	48	-4.512	120.896	-4.155	1.00	54.55
55	ATOM	382	C	GLU	48	0.002	118.641	-0.940	1.00	36.94
	ATOM	383	O	GLU	48	-0.105	119.376	0.033	1.00	40.21
	ATOM	384	N	THR	49	1.171	118.152	-1.331	1.00	36.34
	ATOM	385	CA	THR	49	2.350	118.479	-0.544	1.00	35.48
	ATOM	386	CB	THR	49	3.510	117.473	-0.787	1.00	31.16
60	ATOM	387	OG1	THR	49	4.419	118.011	-1.744	1.00	32.48
	ATOM	388	CG2	THR	49	2.979	116.155	-1.296	1.00	30.36
	ATOM	389	C	THR	49	2.812	119.909	-0.837	1.00	34.59
	ATOM	390	O	THR	49	2.602	120.433	-1.929	1.00	35.13
	ATOM	391	N	GLY	50	3.422	120.537	0.164	1.00	32.49
65	ATOM	392	CA	GLY	50	3.902	121.893	0.007	1.00	28.17
	ATOM	393	C	GLY	50	4.186	122.524	1.351	1.00	29.99

	ATOM	394	O	GLY	50	4.333	121.827	2.342	1.00	29.76
	ATOM	395	N	TYR	51	4.271	123.849	1.380	1.00	31.68
	ATOM	396	CA	TYR	51	4.521	124.584	2.615	1.00	32.88
	ATOM	397	CB	TYR	51	5.547	125.699	2.384	1.00	35.97
5	ATOM	398	CG	TYR	51	6.933	125.177	2.147	1.00	43.03
	ATOM	399	CD1	TYR	51	7.320	124.712	0.887	1.00	44.15
	ATOM	400	CE1	TYR	51	8.577	124.139	0.684	1.00	47.81
	ATOM	401	CD2	TYR	51	7.836	125.066	3.202	1.00	47.08
	ATOM	402	CE2	TYR	51	9.094	124.493	3.016	1.00	51.79
10	ATOM	403	CZ	TYR	51	9.459	124.027	1.755	1.00	51.15
	ATOM	404	OH	TYR	51	10.686	123.413	1.586	1.00	52.20
	ATOM	405	C	TYR	51	3.229	125.178	3.161	1.00	31.97
	ATOM	406	O	TYR	51	2.510	125.884	2.460	1.00	30.01
	ATOM	407	N	PHE	52	2.938	124.886	4.424	1.00	31.50
15	ATOM	408	CA	PHE	52	1.723	125.387	5.043	1.00	30.41
	ATOM	409	CB	PHE	52	0.797	124.229	5.421	1.00	31.30
	ATOM	410	CG	PHE	52	0.379	123.371	4.269	1.00	29.70
	ATOM	411	CD1	PHE	52	1.270	122.487	3.684	1.00	26.20
	ATOM	412	CD2	PHE	52	-0.915	123.441	3.774	1.00	28.38
20	ATOM	413	CE1	PHE	52	0.879	121.690	2.628	1.00	27.17
	ATOM	414	CE2	PHE	52	-1.315	122.646	2.717	1.00	27.76
	ATOM	415	CZ	PHE	52	-0.419	121.770	2.143	1.00	27.27
	ATOM	416	C	PHE	52	1.971	126.207	6.292	1.00	27.65
	ATOM	417	O	PHE	52	2.911	125.959	7.037	1.00	24.85
25	ATOM	418	N	PHE	53	1.113	127.194	6.501	1.00	27.18
	ATOM	419	CA	PHE	53	1.161	128.030	7.690	1.00	27.05
	ATOM	420	CB	PHE	53	0.730	129.456	7.372	1.00	28.76
	ATOM	421	CG	PHE	53	0.572	130.318	8.582	1.00	28.00
	ATOM	422	CD1	PHE	53	1.677	130.736	9.307	1.00	26.49
30	ATOM	423	CD2	PHE	53	-0.690	130.702	9.013	1.00	28.04
	ATOM	424	CE1	PHE	53	1.528	131.525	10.443	1.00	26.54
	ATOM	425	CE2	PHE	53	-0.844	131.490	10.149	1.00	25.78
	ATOM	426	CZ	PHE	53	0.268	131.900	10.862	1.00	24.39
	ATOM	427	C	PHE	53	0.118	127.369	8.587	1.00	27.56
35	ATOM	428	O	PHE	53	-1.044	127.260	8.207	1.00	25.39
	ATOM	429	N	ILE	54	0.538	126.916	9.764	1.00	28.87
	ATOM	430	CA	ILE	54	-0.361	126.232	10.688	1.00	28.95
	ATOM	431	CB	ILE	54	0.189	124.825	11.003	1.00	29.60
	ATOM	432	CG2	ILE	54	-0.865	124.006	11.731	1.00	27.42
40	ATOM	433	CG1	ILE	54	0.583	124.129	9.698	1.00	26.41
	ATOM	434	CD1	ILE	54	1.427	122.904	9.876	1.00	28.70
	ATOM	435	C	ILE	54	-0.538	127.021	11.984	1.00	28.57
	ATOM	436	O	ILE	54	0.421	127.545	12.540	1.00	29.17
	ATOM	437	N	TYR	55	-1.769	127.104	12.473	1.00	26.97
45	ATOM	438	CA	TYR	55	-2.027	127.866	13.690	1.00	24.22
	ATOM	439	CB	TYR	55	-2.490	129.276	13.321	1.00	21.82
	ATOM	440	CG	TYR	55	-3.735	129.317	12.470	1.00	27.68
	ATOM	441	CD1	TYR	55	-4.999	129.382	13.050	1.00	27.01
	ATOM	442	CE1	TYR	55	-6.141	129.398	12.269	1.00	27.44
50	ATOM	443	CD2	TYR	55	-3.652	129.270	11.083	1.00	27.16
	ATOM	444	CE2	TYR	55	-4.792	129.283	10.294	1.00	25.39
	ATOM	445	CZ	TYR	55	-6.029	129.346	10.892	1.00	29.02
	ATOM	446	OH	TYR	55	-7.155	129.347	10.110	1.00	31.82
	ATOM	447	C	TYR	55	-3.040	127.208	14.602	1.00	25.16
55	ATOM	448	O	TYR	55	-3.767	126.332	14.191	1.00	28.47
	ATOM	449	N	GLY	56	-3.075	127.630	15.854	1.00	27.44
	ATOM	450	CA	GLY	56	-4.015	127.050	16.789	1.00	22.07
	ATOM	451	C	GLY	56	-4.088	127.829	18.084	1.00	25.83
	ATOM	452	O	GLY	56	-3.096	128.394	18.531	1.00	26.00
60	ATOM	453	N	GLN	57	-5.277	127.878	18.672	1.00	22.56
	ATOM	454	CA	GLN	57	-5.491	128.566	19.935	1.00	20.91
	ATOM	455	CB	GLN	57	-6.040	129.978	19.726	1.00	17.87
	ATOM	456	CG	GLN	57	-6.397	130.667	21.037	1.00	16.86
	ATOM	457	CD	GLN	57	-6.864	132.098	20.867	1.00	22.06
65	ATOM	458	OE1	GLN	57	-6.108	132.960	20.432	1.00	27.26
	ATOM	459	NE2	GLN	57	-8.118	132.358	21.218	1.00	20.20

	ATOM	460	C	GLN	57	-6.476	127.772	20.767	1.00	24.23
	ATOM	461	O	GLN	57	-7.403	127.169	20.230	1.00	25.16
	ATOM	462	N	VAL	58	-6.260	127.778	22.079	1.00	24.62
	ATOM	463	CA	VAL	58	-7.111	127.074	23.031	1.00	26.68
5	ATOM	464	CB	VAL	58	-6.471	125.739	23.483	1.00	25.55
	ATOM	465	CG1	VAL	58	-7.266	125.137	24.615	1.00	15.69
	ATOM	466	CG2	VAL	58	-6.402	124.767	22.324	1.00	25.77
	ATOM	467	C	VAL	58	-7.258	127.954	24.255	1.00	31.06
	ATOM	468	O	VAL	58	-6.293	128.586	24.663	1.00	27.74
10	ATOM	469	N	LEU	59	-8.458	128.002	24.836	1.00	33.82
	ATOM	470	CA	LEU	59	-8.694	128.785	26.054	1.00	29.85
	ATOM	471	CB	LEU	59	-10.053	129.477	26.018	1.00	29.62
	ATOM	472	CG	LEU	59	-10.236	130.720	26.899	1.00	28.11
	ATOM	473	CD1	LEU	59	-11.712	131.044	27.002	1.00	26.67
15	ATOM	474	CD2	LEU	59	-9.665	130.512	28.271	1.00	22.45
	ATOM	475	C	LEU	59	-8.665	127.823	27.239	1.00	30.98
	ATOM	476	O	LEU	59	-9.521	126.949	27.361	1.00	32.52
	ATOM	477	N	TYR	60	-7.675	127.985	28.110	1.00	33.01
	ATOM	478	CA	TYR	60	-7.549	127.122	29.276	1.00	34.51
20	ATOM	479	CB	TYR	60	-6.077	126.884	29.595	1.00	40.10
	ATOM	480	CG	TYR	60	-5.355	126.187	28.484	1.00	45.37
	ATOM	481	CD1	TYR	60	-4.467	126.876	27.658	1.00	47.13
	ATOM	482	CE1	TYR	60	-3.867	126.246	26.576	1.00	50.28
	ATOM	483	CD2	TYR	60	-5.617	124.852	28.205	1.00	44.35
25	ATOM	484	CE2	TYR	60	-5.029	124.212	27.134	1.00	48.81
	ATOM	485	CZ	TYR	60	-4.160	124.912	26.321	1.00	51.91
	ATOM	486	OH	TYR	60	-3.615	124.276	25.233	1.00	53.77
	ATOM	487	C	TYR	60	-8.252	127.670	30.506	1.00	35.49
	ATOM	488	O	TYR	60	-7.976	128.776	30.959	1.00	36.13
30	ATOM	489	N	THR	61	-9.171	126.878	31.040	1.00	36.19
	ATOM	490	CA	THR	61	-9.913	127.254	32.228	1.00	35.36
	ATOM	491	CB	THR	61	-11.411	127.335	31.932	1.00	34.22
	ATOM	492	OG1	THR	61	-11.843	126.112	31.328	1.00	34.22
	ATOM	493	CG2	THR	61	-11.705	128.493	30.996	1.00	25.45
35	ATOM	494	C	THR	61	-9.644	126.179	33.272	1.00	37.82
	ATOM	495	O	THR	61	-10.494	125.854	34.095	1.00	43.70
	ATOM	496	N	ASP	62	-8.437	125.634	33.211	1.00	37.64
	ATOM	497	CA	ASP	62	-7.974	124.585	34.103	1.00	38.08
	ATOM	498	CB	ASP	62	-7.350	123.486	33.249	1.00	38.42
40	ATOM	499	CG	ASP	62	-6.995	122.259	34.037	1.00	41.25
	ATOM	500	OD1	ASP	62	-7.194	121.142	33.502	1.00	38.28
	ATOM	501	OD2	ASP	62	-6.510	122.415	35.176	1.00	41.37
	ATOM	502	C	ASP	62	-6.937	125.209	35.033	1.00	40.90
	ATOM	503	O	ASP	62	-6.151	126.038	34.595	1.00	46.73
45	ATOM	504	N	LYS	63	-6.921	124.830	36.308	1.00	42.62
	ATOM	505	CA	LYS	63	-5.952	125.430	37.227	1.00	40.92
	ATOM	506	CB	LYS	63	-6.621	125.768	38.560	1.00	42.79
	ATOM	507	CG	LYS	63	-7.242	124.590	39.279	1.00	45.91
	ATOM	508	CD	LYS	63	-8.063	125.070	40.476	1.00	46.22
50	ATOM	509	CE	LYS	63	-9.224	125.976	40.023	1.00	49.97
	ATOM	510	NZ	LYS	63	-9.938	126.665	41.152	1.00	49.98
	ATOM	511	C	LYS	63	-4.693	124.608	37.481	1.00	42.59
	ATOM	512	O	LYS	63	-3.992	124.834	38.460	1.00	42.46
	ATOM	513	N	THR	64	-4.391	123.688	36.574	1.00	44.06
55	ATOM	514	CA	THR	64	-3.224	122.825	36.695	1.00	44.94
	ATOM	515	CB	THR	64	-3.303	121.672	35.663	1.00	43.79
	ATOM	516	OG1	THR	64	-4.542	120.974	35.829	1.00	46.20
	ATOM	517	CG2	THR	64	-2.163	120.679	35.858	1.00	42.80
	ATOM	518	C	THR	64	-1.853	123.509	36.573	1.00	45.82
60	ATOM	519	O	THR	64	-0.970	123.000	35.905	1.00	53.10
	ATOM	520	N	TYR	65	-1.668	124.649	37.221	1.00	45.11
	ATOM	521	CA	TYR	65	-0.390	125.377	37.215	1.00	45.72
	ATOM	522	CB	TYR	65	0.621	124.675	38.141	1.00	40.46
	ATOM	523	CG	TYR	65	1.556	123.691	37.473	1.00	41.90
65	ATOM	524	CD1	TYR	65	2.764	124.110	36.912	1.00	42.47
	ATOM	525	CE1	TYR	65	3.631	123.199	36.282	1.00	41.83

	ATOM	526	CD2	TYR	65	1.231	122.332	37.392	1.00	41.65
	ATOM	527	CE2	TYR	65	2.087	121.412	36.763	1.00	40.28
	ATOM	528	CZ	TYR	65	3.283	121.856	36.212	1.00	41.90
	ATOM	529	OH	TYR	65	4.122	120.962	35.596	1.00	43.84
5	ATOM	530	C	TYR	65	0.270	125.663	35.856	1.00	45.31
	ATOM	531	O	TYR	65	0.965	126.676	35.697	1.00	49.74
	ATOM	532	N	ALA	66	0.068	124.784	34.885	1.00	42.32
	ATOM	533	CA	ALA	66	0.648	124.961	33.560	1.00	39.57
	ATOM	534	CB	ALA	66	2.092	124.510	33.556	1.00	35.36
10	ATOM	535	C	ALA	66	-0.150	124.168	32.541	1.00	39.85
	ATOM	536	O	ALA	66	-0.297	122.950	32.660	1.00	37.60
	ATOM	537	N	MET	67	-0.680	124.867	31.546	1.00	41.88
	ATOM	538	CA	MET	67	-1.462	124.227	30.499	1.00	41.48
	ATOM	539	CB	MET	67	-2.924	124.690	30.546	1.00	39.98
15	ATOM	540	CG	MET	67	-3.709	124.211	31.760	1.00	37.31
	ATOM	541	SD	MET	67	-3.776	122.420	31.895	1.00	35.46
	ATOM	542	CE	MET	67	-5.089	122.015	30.741	1.00	34.06
	ATOM	543	C	MET	67	-0.858	124.566	29.147	1.00	39.50
	ATOM	544	O	MET	67	-0.024	125.467	29.035	1.00	38.44
20	ATOM	545	N	GLY	68	-1.280	123.835	28.123	1.00	39.84
	ATOM	546	CA	GLY	68	-0.769	124.076	26.789	1.00	37.61
	ATOM	547	C	GLY	68	-1.187	122.999	25.818	1.00	33.76
	ATOM	548	O	GLY	68	-1.768	121.995	26.213	1.00	35.58
	ATOM	549	N	HIS	69	-0.905	123.211	24.540	1.00	35.07
25	ATOM	550	CA	HIS	69	-1.252	122.235	23.521	1.00	33.70
	ATOM	551	CB	HIS	69	-2.529	122.655	22.768	1.00	37.02
	ATOM	552	CG	HIS	69	-2.485	124.038	22.194	1.00	34.65
	ATOM	553	CD2	HIS	69	-2.341	124.467	20.920	1.00	35.63
	ATOM	554	ND1	HIS	69	-2.617	125.172	22.965	1.00	37.68
30	ATOM	555	CE1	HIS	69	-2.556	126.241	22.192	1.00	32.64
	ATOM	556	NE2	HIS	69	-2.390	125.840	20.945	1.00	35.90
	ATOM	557	C	HIS	69	-0.111	122.024	22.536	1.00	33.98
	ATOM	558	O	HIS	69	0.861	122.762	22.536	1.00	30.94
	ATOM	559	N	LEU	70	-0.237	121.003	21.701	1.00	35.35
35	ATOM	560	CA	LEU	70	0.780	120.689	20.709	1.00	30.76
	ATOM	561	CB	LEU	70	1.423	119.341	21.017	1.00	29.59
	ATOM	562	CG	LEU	70	1.723	118.976	22.468	1.00	30.71
	ATOM	563	CD1	LEU	70	2.106	117.518	22.542	1.00	25.60
	ATOM	564	CD2	LEU	70	2.825	119.859	23.005	1.00	27.39
40	ATOM	565	C	LEU	70	0.151	120.585	19.323	1.00	33.19
	ATOM	566	O	LEU	70	-0.934	120.025	19.169	1.00	37.07
	ATOM	567	N	ILE	71	0.811	121.145	18.317	1.00	30.93
	ATOM	568	CA	ILE	71	0.317	121.012	16.956	1.00	27.94
	ATOM	569	CB	ILE	71	0.485	122.299	16.144	1.00	28.22
45	ATOM	570	CG2	ILE	71	0.284	122.017	14.670	1.00	24.54
	ATOM	571	CG1	ILE	71	-0.536	123.330	16.608	1.00	26.34
	ATOM	572	CD1	ILE	71	-0.329	124.695	16.014	1.00	37.50
	ATOM	573	C	ILE	71	1.245	119.930	16.428	1.00	30.32
	ATOM	574	O	ILE	71	2.449	120.141	16.341	1.00	23.07
50	ATOM	575	N	GLN	72	0.693	118.763	16.108	1.00	29.74
	ATOM	576	CA	GLN	72	1.527	117.664	15.645	1.00	27.85
	ATOM	577	CB	GLN	72	1.332	116.468	16.564	1.00	26.70
	ATOM	578	CG	GLN	72	1.474	116.829	18.015	1.00	28.28
	ATOM	579	CD	GLN	72	1.479	115.621	18.907	1.00	33.11
55	ATOM	580	OE1	GLN	72	0.571	114.797	18.851	1.00	41.22
	ATOM	581	NE2	GLN	72	2.501	115.506	19.744	1.00	30.18
	ATOM	582	C	GLN	72	1.315	117.234	14.208	1.00	29.85
	ATOM	583	O	GLN	72	0.260	117.473	13.618	1.00	28.20
	ATOM	584	N	ARG	73	2.334	116.579	13.664	1.00	30.90
60	ATOM	585	CA	ARG	73	2.301	116.087	12.296	1.00	30.79
	ATOM	586	CB	ARG	73	3.362	116.807	11.475	1.00	29.18
	ATOM	587	CG	ARG	73	3.478	116.314	10.064	1.00	28.66
	ATOM	588	CD	ARG	73	4.837	116.631	9.517	1.00	29.25
	ATOM	589	NE	ARG	73	4.969	116.212	8.130	1.00	35.78
65	ATOM	590	CZ	ARG	73	6.127	116.045	7.510	1.00	35.27
	ATOM	591	NH1	ARG	73	7.264	116.255	8.158	1.00	34.95

	ATOM	592	NH2	ARG	73	6.145	115.673	6.243	1.00	33.33
	ATOM	593	C	ARG	73	2.552	114.583	12.226	1.00	32.42
	ATOM	594	O	ARG	73	3.524	114.089	12.787	1.00	34.19
	ATOM	595	N	LYS	74	1.660	113.860	11.555	1.00	35.90
5	ATOM	596	CA	LYS	74	1.819	112.418	11.371	1.00	38.57
	ATOM	597	CB	LYS	74	0.483	111.685	11.523	1.00	43.52
	ATOM	598	CG	LYS	74	-0.034	111.590	12.949	1.00	52.89
	ATOM	599	CD	LYS	74	-1.406	110.882	13.028	1.00	58.14
	ATOM	600	CE	LYS	74	-1.906	110.816	14.472	1.00	56.75
10	ATOM	601	NZ	LYS	74	-3.251	110.176	14.577	1.00	63.98
	ATOM	602	C	LYS	74	2.329	112.201	9.952	1.00	40.11
	ATOM	603	O	LYS	74	1.570	112.345	8.985	1.00	37.55
	ATOM	604	N	LYS	75	3.606	111.857	9.827	1.00	34.48
	ATOM	605	CA	LYS	75	4.215	111.633	8.521	1.00	34.24
15	ATOM	606	CB	LYS	75	5.728	111.457	8.674	1.00	36.16
	ATOM	607	CG	LYS	75	6.483	112.599	9.349	1.00	35.99
	ATOM	608	CD	LYS	75	7.975	112.248	9.432	1.00	40.91
	ATOM	609	CE	LYS	75	8.792	113.361	10.074	1.00	46.16
	ATOM	610	NZ	LYS	75	10.256	113.051	10.234	1.00	46.03
20	ATOM	611	C	LYS	75	3.643	110.386	7.839	1.00	30.92
	ATOM	612	O	LYS	75	3.418	109.382	8.488	1.00	27.47
	ATOM	613	N	VAL	76	3.417	110.459	6.529	1.00	32.82
	ATOM	614	CA	VAL	76	2.906	109.318	5.765	1.00	31.51
	ATOM	615	CB	VAL	76	2.367	109.713	4.378	1.00	29.19
25	ATOM	616	CG1	VAL	76	1.195	108.844	4.023	1.00	27.09
	ATOM	617	CG2	VAL	76	2.005	111.155	4.344	1.00	34.94
	ATOM	618	C	VAL	76	4.075	108.395	5.488	1.00	33.40
	ATOM	619	O	VAL	76	3.950	107.177	5.548	1.00	32.92
	ATOM	620	N	HIS	77	5.211	109.003	5.160	1.00	34.88
30	ATOM	621	CA	HIS	77	6.430	108.278	4.855	1.00	34.59
	ATOM	622	CB	HIS	77	7.055	108.819	3.565	1.00	35.92
	ATOM	623	CG	HIS	77	6.147	108.746	2.376	1.00	34.17
	ATOM	624	CD2	HIS	77	6.113	109.473	1.236	1.00	31.44
	ATOM	625	ND1	HIS	77	5.139	107.810	2.262	1.00	35.05
35	ATOM	626	CE1	HIS	77	4.522	107.964	1.106	1.00	29.84
	ATOM	627	NE2	HIS	77	5.095	108.964	0.464	1.00	38.35
	ATOM	628	C	HIS	77	7.399	108.421	6.018	1.00	35.63
	ATOM	629	O	HIS	77	7.530	109.504	6.588	1.00	40.24
	ATOM	630	N	VAL	78	8.096	107.337	6.352	1.00	37.37
40	ATOM	631	CA	VAL	78	9.001	107.371	7.487	1.00	41.32
	ATOM	632	CB	VAL	78	8.512	106.392	8.572	1.00	43.10
	ATOM	633	CG1	VAL	78	9.362	106.525	9.811	1.00	45.01
	ATOM	634	CG2	VAL	78	7.069	106.703	8.930	1.00	44.98
	ATOM	635	C	VAL	78	10.511	107.196	7.280	1.00	43.02
45	ATOM	636	O	VAL	78	11.261	108.142	7.513	1.00	49.79
	ATOM	637	N	PHE	79	10.986	106.028	6.869	1.00	41.33
	ATOM	638	CA	PHE	79	12.447	105.844	6.686	1.00	45.00
	ATOM	639	CB	PHE	79	13.125	107.056	6.016	1.00	40.51
	ATOM	640	CG	PHE	79	12.530	107.454	4.708	1.00	39.52
50	ATOM	641	CD1	PHE	79	11.729	108.586	4.617	1.00	36.80
	ATOM	642	CD2	PHE	79	12.772	106.703	3.565	1.00	37.89
	ATOM	643	CE1	PHE	79	11.177	108.968	3.405	1.00	37.52
	ATOM	644	CE2	PHE	79	12.224	107.078	2.349	1.00	35.59
	ATOM	645	CZ	PHE	79	11.424	108.213	2.267	1.00	37.58
55	ATOM	646	C	PHE	79	13.225	105.614	7.982	1.00	43.50
	ATOM	647	O	PHE	79	13.179	106.428	8.906	1.00	37.85
	ATOM	648	N	GLY	80	13.971	104.515	8.017	1.00	46.65
	ATOM	649	CA	GLY	80	14.794	104.188	9.164	1.00	48.38
	ATOM	650	C	GLY	80	14.166	104.312	10.535	1.00	47.89
60	ATOM	651	O	GLY	80	13.060	103.819	10.774	1.00	51.15
	ATOM	652	N	ASP	81	14.880	104.978	11.440	1.00	45.07
	ATOM	653	CA	ASP	81	14.415	105.145	12.804	1.00	43.99
	ATOM	654	CB	ASP	81	15.585	104.979	13.782	1.00	45.66
	ATOM	655	CG	ASP	81	16.589	106.118	13.704	1.00	47.49
65	ATOM	656	OD1	ASP	81	16.549	106.907	12.734	1.00	51.90
	ATOM	657	OD2	ASP	81	17.433	106.222	14.616	1.00	48.08

	ATOM	658	C	ASP	81	13.703	106.461	13.064	1.00	42.29
	ATOM	659	O	ASP	81	13.642	106.918	14.205	1.00	39.94
	ATOM	660	N	GLU	82	13.167	107.071	12.012	1.00	39.91
	ATOM	661	CA	GLU	82	12.431	108.324	12.165	1.00	36.58
5	ATOM	662	CB	GLU	82	11.956	108.872	10.818	1.00	34.91
	ATOM	663	CG	GLU	82	12.958	109.591	9.974	1.00	37.82
	ATOM	664	CD	GLU	82	12.281	110.563	9.022	1.00	40.75
	ATOM	665	OE1	GLU	82	11.256	110.194	8.415	1.00	38.15
	ATOM	666	OE2	GLU	82	12.768	111.704	8.878	1.00	47.99
10	ATOM	667	C	GLU	82	11.175	108.070	12.987	1.00	36.58
	ATOM	668	O	GLU	82	10.601	106.985	12.936	1.00	38.40
	ATOM	669	N	LEU	83	10.751	109.073	13.743	1.00	33.83
	ATOM	670	CA	LEU	83	9.519	108.978	14.509	1.00	32.70
	ATOM	671	CB	LEU	83	9.567	109.893	15.732	1.00	30.78
15	ATOM	672	CG	LEU	83	10.051	109.299	17.053	1.00	29.26
	ATOM	673	CD1	LEU	83	11.260	108.424	16.839	1.00	29.60
	ATOM	674	CD2	LEU	83	10.366	110.422	18.008	1.00	28.65
	ATOM	675	C	LEU	83	8.504	109.501	13.515	1.00	33.88
	ATOM	676	O	LEU	83	8.739	110.528	12.887	1.00	38.46
20	ATOM	677	N	SER	84	7.389	108.804	13.351	1.00	34.94
	ATOM	678	CA	SER	84	6.390	109.248	12.391	1.00	39.77
	ATOM	679	CB	SER	84	5.529	108.071	11.936	1.00	38.92
	ATOM	680	OG	SER	84	4.890	107.474	13.038	1.00	43.11
	ATOM	681	C	SER	84	5.502	110.371	12.919	1.00	39.89
25	ATOM	682	O	SER	84	4.753	110.993	12.167	1.00	42.52
	ATOM	683	N	LEU	85	5.594	110.631	14.215	1.00	40.92
	ATOM	684	CA	LEU	85	4.813	111.685	14.832	1.00	34.66
	ATOM	685	CB	LEU	85	4.061	111.135	16.044	1.00	35.12
	ATOM	686	CG	LEU	85	2.950	111.940	16.734	1.00	36.92
30	ATOM	687	CD1	LEU	85	3.530	113.151	17.423	1.00	38.24
	ATOM	688	CD2	LEU	85	1.899	112.346	15.711	1.00	36.08
	ATOM	689	C	LEU	85	5.783	112.773	15.253	1.00	35.30
	ATOM	690	O	LEU	85	6.645	112.563	16.101	1.00	32.85
	ATOM	691	N	VAL	86	5.654	113.936	14.632	1.00	33.26
35	ATOM	692	CA	VAL	86	6.517	115.063	14.944	1.00	34.57
	ATOM	693	CB	VAL	86	7.220	115.624	13.674	1.00	35.39
	ATOM	694	CG1	VAL	86	7.971	116.898	14.007	1.00	31.44
	ATOM	695	CG2	VAL	86	8.178	114.592	13.110	1.00	38.08
	ATOM	696	C	VAL	86	5.682	116.174	15.534	1.00	32.85
40	ATOM	697	O	VAL	86	4.619	116.501	15.013	1.00	35.10
	ATOM	698	N	THR	87	6.142	116.754	16.629	1.00	33.41
	ATOM	699	CA	THR	87	5.398	117.854	17.193	1.00	33.56
	ATOM	700	CB	THR	87	5.215	117.697	18.735	1.00	32.70
	ATOM	701	OG1	THR	87	5.381	118.964	19.384	1.00	34.29
45	ATOM	702	CG2	THR	87	6.176	116.678	19.288	1.00	34.95
	ATOM	703	C	THR	87	6.065	119.185	16.822	1.00	30.57
	ATOM	704	O	THR	87	7.205	119.455	17.190	1.00	28.08
	ATOM	705	N	LEU	88	5.348	119.970	16.024	1.00	28.42
	ATOM	706	CA	LEU	88	5.792	121.285	15.600	1.00	31.31
50	ATOM	707	CB	LEU	88	5.220	121.643	14.221	1.00	28.21
	ATOM	708	CG	LEU	88	5.275	120.856	12.907	1.00	29.12
	ATOM	709	CD1	LEU	88	6.168	119.671	13.017	1.00	31.12
	ATOM	710	CD2	LEU	88	3.875	120.441	12.525	1.00	28.72
	ATOM	711	C	LEU	88	5.184	122.274	16.609	1.00	41.15
55	ATOM	712	O	LEU	88	4.099	122.045	17.150	1.00	52.30
	ATOM	713	N	PHE	89	5.869	123.364	16.897	1.00	39.23
	ATOM	714	CA	PHE	89	5.275	124.367	17.784	1.00	42.27
	ATOM	715	CB	PHE	89	4.035	124.924	17.079	1.00	35.63
	ATOM	716	CG	PHE	89	4.261	125.147	15.607	1.00	35.06
60	ATOM	717	CD1	PHE	89	3.302	124.769	14.673	1.00	31.56
	ATOM	718	CD2	PHE	89	5.501	125.625	15.148	1.00	30.96
	ATOM	719	CE1	PHE	89	3.577	124.846	13.304	1.00	33.76
	ATOM	720	CE2	PHE	89	5.785	125.704	13.790	1.00	27.38
	ATOM	721	CZ	PHE	89	4.825	125.311	12.865	1.00	30.29
65	ATOM	722	C	PHE	89	4.983	123.996	19.255	1.00	39.99
	ATOM	723	O	PHE	89	5.916	123.709	20.012	1.00	45.34

	ATOM	724	N	ARG	90	3.733	124.018	19.692	1.00	38.92
	ATOM	725	CA	ARG	90	3.461	123.727	21.112	1.00	41.66
	ATOM	726	CB	ARG	90	4.318	122.543	21.607	1.00	39.24
	ATOM	727	CG	ARG	90	4.825	122.669	23.058	1.00	29.41
5	ATOM	728	CD	ARG	90	5.648	121.445	23.500	1.00	31.13
	ATOM	729	NE	ARG	90	6.357	121.677	24.762	1.00	31.75
	ATOM	730	CZ	ARG	90	7.684	121.743	24.894	1.00	28.86
	ATOM	731	NH1	ARG	90	8.482	121.587	23.851	1.00	25.48
	ATOM	732	NH2	ARG	90	8.218	121.999	26.077	1.00	29.84
10	ATOM	733	C	ARG	90	3.709	124.914	22.063	1.00	39.23
	ATOM	734	O	ARG	90	4.770	125.534	22.033	1.00	31.33
	ATOM	735	N	CYS	91	2.721	125.223	22.907	1.00	40.12
	ATOM	736	CA	CYS	91	2.878	126.295	23.878	1.00	37.09
	ATOM	737	C	CYS	91	2.397	126.013	25.272	1.00	37.21
15	ATOM	738	O	CYS	91	1.693	125.041	25.515	1.00	37.46
	ATOM	739	CB	CYS	91	2.249	127.588	23.412	1.00	40.13
	ATOM	740	SG	CYS	91	0.460	127.750	23.072	1.00	41.64
	ATOM	741	N	ILE	92	2.790	126.892	26.188	1.00	34.02
	ATOM	742	CA	ILE	92	2.452	126.748	27.596	1.00	33.22
20	ATOM	743	CB	ILE	92	3.675	126.239	28.386	1.00	32.88
	ATOM	744	CG2	ILE	92	3.277	125.905	29.807	1.00	32.61
	ATOM	745	CG1	ILE	92	4.249	124.996	27.702	1.00	34.64
	ATOM	746	CD1	ILE	92	5.572	124.517	28.276	1.00	32.30
	ATOM	747	C	ILE	92	1.984	128.052	28.231	1.00	30.78
25	ATOM	748	O	ILE	92	2.345	129.122	27.788	1.00	33.43
	ATOM	749	N	GLN	93	1.152	127.947	29.260	1.00	29.35
	ATOM	750	CA	GLN	93	0.659	129.104	29.988	1.00	28.69
	ATOM	751	CB	GLN	93	-0.733	129.527	29.516	1.00	28.17
	ATOM	752	CG	GLN	93	-0.810	130.438	28.288	1.00	25.87
30	ATOM	753	CD	GLN	93	-0.114	131.769	28.468	1.00	24.22
	ATOM	754	OE1	GLN	93	1.070	131.895	28.198	1.00	28.90
	ATOM	755	NE2	GLN	93	-0.851	132.771	28.929	1.00	25.29
	ATOM	756	C	GLN	93	0.555	128.706	31.443	1.00	31.44
	ATOM	757	O	GLN	93	-0.063	127.701	31.759	1.00	32.07
35	ATOM	758	N	ASN	94	1.181	129.465	32.331	1.00	35.77
	ATOM	759	CA	ASN	94	1.068	129.165	33.746	1.00	33.56
	ATOM	760	CB	ASN	94	2.054	129.999	34.565	1.00	34.70
	ATOM	761	CG	ASN	94	3.444	129.415	34.575	1.00	33.01
	ATOM	762	OD1	ASN	94	3.628	128.248	34.876	1.00	30.56
40	ATOM	763	ND2	ASN	94	4.432	130.230	34.263	1.00	36.61
	ATOM	764	C	ASN	94	-0.369	129.533	34.115	1.00	35.80
	ATOM	765	O	ASN	94	-0.922	130.506	33.598	1.00	36.63
	ATOM	766	N	MET	95	-0.981	128.754	34.997	1.00	35.73
	ATOM	767	CA	MET	95	-2.357	129.017	35.401	1.00	34.73
45	ATOM	768	CB	MET	95	-3.197	127.750	35.217	1.00	32.97
	ATOM	769	CG	MET	95	-3.145	127.139	33.824	1.00	27.35
	ATOM	770	SD	MET	95	-3.583	128.288	32.503	1.00	25.14
	ATOM	771	CE	MET	95	-5.324	128.466	32.702	1.00	15.55
	ATOM	772	C	MET	95	-2.462	129.496	36.850	1.00	36.36
50	ATOM	773	O	MET	95	-1.610	129.172	37.682	1.00	40.70
	ATOM	774	N	PRO	96	-3.493	130.299	37.161	1.00	36.06
	ATOM	775	CD	PRO	96	-4.314	131.036	36.201	1.00	35.32
	ATOM	776	CA	PRO	96	-3.724	130.817	38.510	1.00	40.00
	ATOM	777	CB	PRO	96	-4.532	132.089	38.272	1.00	33.97
55	ATOM	778	CG	PRO	96	-4.346	132.383	36.838	1.00	37.05
	ATOM	779	C	PRO	96	-4.557	129.776	39.253	1.00	46.34
	ATOM	780	O	PRO	96	-5.091	128.855	38.637	1.00	47.59
	ATOM	781	N	GLU	97	-4.689	129.924	40.567	1.00	52.14
	ATOM	782	CA	GLU	97	-5.469	128.973	41.351	1.00	57.27
60	ATOM	783	CB	GLU	97	-4.976	128.944	42.798	1.00	62.29
	ATOM	784	CG	GLU	97	-5.298	127.657	43.551	1.00	72.31
	ATOM	785	CD	GLU	97	-4.097	126.700	43.606	1.00	79.34
	ATOM	786	OE1	GLU	97	-3.042	127.087	44.199	1.00	84.94
	ATOM	787	OE2	GLU	97	-4.207	125.569	43.056	1.00	81.39
65	ATOM	788	C	GLU	97	-6.934	129.394	41.332	1.00	56.67
	ATOM	789	O	GLU	97	-7.836	128.553	41.297	1.00	59.18

	ATOM	790	N	THR	98	-7.162	130.703	41.334	1.00	57.55
	ATOM	791	CA	THR	98	-8.511	131.256	41.361	1.00	59.71
	ATOM	792	CB	THR	98	-8.466	132.754	41.632	1.00	59.24
	ATOM	793	OG1	THR	98	-7.683	133.382	40.599	1.00	68.81
5	ATOM	794	CG2	THR	98	-7.855	133.024	43.023	1.00	56.15
	ATOM	795	C	THR	98	-9.381	131.036	40.125	1.00	60.32
	ATOM	796	O	THR	98	-10.027	129.988	40.001	1.00	65.03
	ATOM	797	N	LEU	99	-9.420	132.022	39.227	1.00	54.94
	ATOM	798	CA	LEU	99	-10.248	131.928	38.022	1.00	51.13
10	ATOM	799	CB	LEU	99	-11.088	133.195	37.880	1.00	50.56
	ATOM	800	CG	LEU	99	-12.255	133.354	38.851	1.00	52.80
	ATOM	801	CD1	LEU	99	-12.718	134.809	38.889	1.00	48.38
	ATOM	802	CD2	LEU	99	-13.385	132.430	38.422	1.00	50.80
	ATOM	803	C	LEU	99	-9.453	131.695	36.733	1.00	50.20
15	ATOM	804	O	LEU	99	-9.249	132.623	35.937	1.00	50.50
	ATOM	805	N	PRO	100	-9.024	130.443	36.492	1.00	47.97
	ATOM	806	CD	PRO	100	-9.364	129.227	37.246	1.00	45.68
	ATOM	807	CA	PRO	100	-8.246	130.102	35.298	1.00	46.54
	ATOM	808	CB	PRO	100	-8.132	128.579	35.377	1.00	45.50
20	ATOM	809	CG	PRO	100	-8.259	128.296	36.838	1.00	45.80
	ATOM	810	C	PRO	100	-8.913	130.550	34.009	1.00	46.78
	ATOM	811	O	PRO	100	-10.046	130.180	33.724	1.00	52.13
	ATOM	812	N	ASN	101	-8.204	131.351	33.231	1.00	42.71
	ATOM	813	CA	ASN	101	-8.705	131.830	31.958	1.00	41.37
25	ATOM	814	CB	ASN	101	-9.753	132.915	32.171	1.00	43.24
	ATOM	815	CG	ASN	101	-11.136	132.349	32.425	1.00	45.93
	ATOM	816	OD1	ASN	101	-11.765	131.786	31.526	1.00	45.33
	ATOM	817	ND2	ASN	101	-11.618	132.490	33.658	1.00	48.11
	ATOM	818	C	ASN	101	-7.534	132.385	31.171	1.00	42.66
30	ATOM	819	O	ASN	101	-7.295	133.595	31.163	1.00	44.79
	ATOM	820	N	ASN	102	-6.785	131.507	30.510	1.00	40.46
	ATOM	821	CA	ASN	102	-5.655	131.991	29.752	1.00	37.90
	ATOM	822	CB	ASN	102	-4.367	131.360	30.256	1.00	34.01
	ATOM	823	CG	ASN	102	-3.766	132.138	31.410	1.00	32.44
35	ATOM	824	OD1	ASN	102	-3.768	133.366	31.404	1.00	29.07
	ATOM	825	ND2	ASN	102	-3.244	131.429	32.400	1.00	33.79
	ATOM	826	C	ASN	102	-5.702	131.935	28.238	1.00	39.20
	ATOM	827	O	ASN	102	-5.693	132.977	27.594	1.00	46.07
	ATOM	828	N	SER	103	-5.759	130.761	27.640	1.00	35.42
40	ATOM	829	CA	SER	103	-5.762	130.721	26.167	1.00	40.28
	ATOM	830	CB	SER	103	-6.872	131.620	25.569	1.00	40.68
	ATOM	831	OG	SER	103	-6.374	132.857	25.071	1.00	31.76
	ATOM	832	C	SER	103	-4.384	131.150	25.607	1.00	35.24
	ATOM	833	O	SER	103	-3.881	132.239	25.877	1.00	24.65
45	ATOM	834	N	CYS	104	-3.776	130.249	24.847	1.00	32.88
	ATOM	835	CA	CYS	104	-2.486	130.491	24.256	1.00	31.77
	ATOM	836	C	CYS	104	-2.601	130.244	22.742	1.00	30.58
	ATOM	837	O	CYS	104	-3.210	129.273	22.304	1.00	29.23
	ATOM	838	CB	CYS	104	-1.411	129.569	24.895	1.00	28.43
50	ATOM	839	SG	CYS	104	0.065	129.638	23.867	1.00	46.07
	ATOM	840	N	TYR	105	-2.040	131.153	21.945	1.00	30.34
	ATOM	841	CA	TYR	105	-2.035	131.029	20.489	1.00	26.12
	ATOM	842	CB	TYR	105	-2.521	132.323	19.837	1.00	27.03
	ATOM	843	CG	TYR	105	-2.374	132.371	18.324	1.00	25.61
55	ATOM	844	CD1	TYR	105	-3.449	132.074	17.479	1.00	24.90
	ATOM	845	CE1	TYR	105	-3.314	132.116	16.099	1.00	26.08
	ATOM	846	CD2	TYR	105	-1.160	132.709	17.735	1.00	25.41
	ATOM	847	CE2	TYR	105	-1.018	132.749	16.354	1.00	28.91
	ATOM	848	CZ	TYR	105	-2.098	132.453	15.543	1.00	27.58
60	ATOM	849	OH	TYR	105	-1.956	132.508	14.174	1.00	29.89
	ATOM	850	C	TYR	105	-0.602	130.757	20.038	1.00	29.20
	ATOM	851	O	TYR	105	0.348	131.278	20.614	1.00	33.10
	ATOM	852	N	SER	106	-0.441	129.940	19.011	1.00	30.48
	ATOM	853	CA	SER	106	0.891	129.639	18.512	1.00	29.22
65	ATOM	854	CB	SER	106	1.527	128.528	19.352	1.00	28.30
	ATOM	855	OG	SER	106	2.845	128.242	18.929	1.00	28.62

	ATOM	856	C	SER	106	0.763	129.203	17.063	1.00	29.94
	ATOM	857	O	SER	106	-0.201	128.538	16.706	1.00	29.72
	ATOM	858	N	ALA	107	1.721	129.597	16.228	1.00	27.64
	ATOM	859	CA	ALA	107	1.699	129.235	14.817	1.00	24.33
5	ATOM	860	CB	ALA	107	0.831	130.209	14.040	1.00	21.30
	ATOM	861	C	ALA	107	3.097	129.199	14.220	1.00	25.56
	ATOM	862	O	ALA	107	4.051	129.696	14.804	1.00	28.03
	ATOM	863	N	GLY	108	3.210	128.602	13.044	1.00	26.07
	ATOM	864	CA	GLY	108	4.493	128.522	12.377	1.00	25.49
10	ATOM	865	C	GLY	108	4.337	127.880	11.018	1.00	29.03
	ATOM	866	O	GLY	108	3.249	127.452	10.661	1.00	32.20
	ATOM	867	N	ILE	109	5.420	127.811	10.257	1.00	29.34
	ATOM	868	CA	ILE	109	5.384	127.213	8.931	1.00	27.71
	ATOM	869	CB	ILE	109	6.078	128.120	7.901	1.00	27.48
15	ATOM	870	CG2	ILE	109	6.052	127.475	6.531	1.00	26.46
	ATOM	871	CG1	ILE	109	5.377	129.477	7.864	1.00	26.16
	ATOM	872	CD1	ILE	109	6.071	130.502	7.013	1.00	29.07
	ATOM	873	C	ILE	109	6.083	125.868	8.956	1.00	28.66
	ATOM	874	O	ILE	109	7.086	125.693	9.646	1.00	30.59
20	ATOM	875	N	ALA	110	5.551	124.910	8.209	1.00	25.60
	ATOM	876	CA	ALA	110	6.136	123.576	8.146	1.00	26.36
	ATOM	877	CB	ALA	110	5.521	122.690	9.203	1.00	20.26
	ATOM	878	C	ALA	110	5.899	122.972	6.773	1.00	29.61
	ATOM	879	O	ALA	110	4.960	123.341	6.084	1.00	27.61
25	ATOM	880	N	LYS	111	6.762	122.056	6.359	1.00	33.77
	ATOM	881	CA	LYS	111	6.566	121.421	5.077	1.00	35.09
	ATOM	882	CB	LYS	111	7.891	121.153	4.374	1.00	39.38
	ATOM	883	CG	LYS	111	7.678	120.339	3.104	1.00	49.74
	ATOM	884	CD	LYS	111	8.637	120.699	2.002	1.00	53.95
30	ATOM	885	CE	LYS	111	8.252	119.974	0.720	1.00	54.28
	ATOM	886	NZ	LYS	111	9.214	120.299	-0.386	1.00	60.85
	ATOM	887	C	LYS	111	5.821	120.112	5.308	1.00	35.28
	ATOM	888	O	LYS	111	6.238	119.284	6.110	1.00	36.58
	ATOM	889	N	LEU	112	4.709	119.943	4.605	1.00	32.32
35	ATOM	890	CA	LEU	112	3.887	118.756	4.736	1.00	29.67
	ATOM	891	CB	LEU	112	2.493	119.143	5.223	1.00	27.04
	ATOM	892	CG	LEU	112	2.434	120.057	6.446	1.00	24.38
	ATOM	893	CD1	LEU	112	0.999	120.460	6.717	1.00	27.56
	ATOM	894	CD2	LEU	112	3.028	119.356	7.637	1.00	19.52
40	ATOM	895	C	LEU	112	3.777	118.043	3.397	1.00	32.58
	ATOM	896	O	LEU	112	4.099	118.611	2.354	1.00	31.51
	ATOM	897	N	GLU	113	3.313	116.800	3.428	1.00	35.75
	ATOM	898	CA	GLU	113	3.164	116.016	2.215	1.00	40.08
	ATOM	899	CB	GLU	113	4.212	114.920	2.146	1.00	44.23
45	ATOM	900	CG	GLU	113	5.619	115.343	2.447	1.00	50.12
	ATOM	901	CD	GLU	113	6.607	114.359	1.863	1.00	58.08
	ATOM	902	OE1	GLU	113	6.318	113.127	1.900	1.00	61.13
	ATOM	903	OE2	GLU	113	7.667	114.821	1.366	1.00	63.64
	ATOM	904	C	GLU	113	1.817	115.341	2.140	1.00	41.63
50	ATOM	905	O	GLU	113	1.146	115.167	3.148	1.00	48.29
	ATOM	906	N	GLU	114	1.440	114.951	0.927	1.00	40.97
	ATOM	907	CA	GLU	114	0.190	114.244	0.681	1.00	36.68
	ATOM	908	CB	GLU	114	0.205	113.620	-0.698	1.00	41.19
	ATOM	909	CG	GLU	114	-0.452	114.390	-1.787	1.00	46.58
55	ATOM	910	CD	GLU	114	-0.929	113.455	-2.877	1.00	51.12
	ATOM	911	OE1	GLU	114	-1.854	112.654	-2.598	1.00	51.88
	ATOM	912	OE2	GLU	114	-0.370	113.510	-4.000	1.00	57.08
	ATOM	913	C	GLU	114	0.079	113.097	1.664	1.00	35.51
	ATOM	914	O	GLU	114	0.953	112.240	1.711	1.00	38.92
60	ATOM	915	N	GLY	115	-0.995	113.065	2.437	1.00	32.99
	ATOM	916	CA	GLY	115	-1.154	111.972	3.370	1.00	31.79
	ATOM	917	C	GLY	115	-0.804	112.321	4.791	1.00	34.39
	ATOM	918	O	GLY	115	-1.177	111.598	5.705	1.00	39.20
	ATOM	919	N	ASP	116	-0.067	113.407	4.984	1.00	35.66
65	ATOM	920	CA	ASP	116	0.284	113.827	6.330	1.00	38.09
	ATOM	921	CB	ASP	116	1.271	115.001	6.308	1.00	36.00

	ATOM	922	CG	ASP	116	2.689	114.578	5.991	1.00	35.09
	ATOM	923	OD1	ASP	116	2.996	113.371	6.070	1.00	33.81
	ATOM	924	OD2	ASP	116	3.506	115.463	5.679	1.00	35.24
	ATOM	925	C	ASP	116	-0.997	114.278	7.015	1.00	39.38
5	ATOM	926	O	ASP	116	-1.948	114.707	6.355	1.00	37.91
	ATOM	927	N	GLU	117	-1.032	114.168	8.337	1.00	41.54
	ATOM	928	CA	GLU	117	-2.198	114.611	9.089	1.00	37.92
	ATOM	929	CB	GLU	117	-2.921	113.428	9.722	1.00	38.90
	ATOM	930	CG	GLU	117	-3.382	112.389	8.732	1.00	48.15
10	ATOM	931	CD	GLU	117	-4.130	111.256	9.401	1.00	52.62
	ATOM	932	OE1	GLU	117	-3.734	110.855	10.530	1.00	55.49
	ATOM	933	OE2	GLU	117	-5.109	110.761	8.789	1.00	57.10
	ATOM	934	C	GLU	117	-1.724	115.550	10.180	1.00	34.02
	ATOM	935	O	GLU	117	-0.641	115.366	10.734	1.00	33.31
15	ATOM	936	N	LEU	118	-2.519	116.572	10.465	1.00	30.68
	ATOM	937	CA	LEU	118	-2.193	117.524	11.513	1.00	27.14
	ATOM	938	CB	LEU	118	-2.286	118.957	10.986	1.00	26.49
	ATOM	939	CG	LEU	118	-1.299	119.401	9.913	1.00	26.21
	ATOM	940	CD1	LEU	118	-1.628	120.812	9.484	1.00	23.05
20	ATOM	941	CD2	LEU	118	0.116	119.323	10.447	1.00	25.38
	ATOM	942	C	LEU	118	-3.197	117.350	12.646	1.00	28.33
	ATOM	943	O	LEU	118	-4.366	117.046	12.403	1.00	31.46
	ATOM	944	N	GLN	119	-2.748	117.526	13.884	1.00	28.96
	ATOM	945	CA	GLN	119	-3.653	117.420	15.021	1.00	29.13
25	ATOM	946	CB	GLN	119	-3.735	115.977	15.498	1.00	28.23
	ATOM	947	CG	GLN	119	-2.485	115.476	16.155	1.00	36.38
	ATOM	948	CD	GLN	119	-2.590	114.022	16.549	1.00	36.57
	ATOM	949	OE1	GLN	119	-1.788	113.531	17.342	1.00	40.70
	ATOM	950	NE2	GLN	119	-3.571	113.322	15.993	1.00	36.80
30	ATOM	951	C	GLN	119	-3.237	118.333	16.170	1.00	29.43
	ATOM	952	O	GLN	119	-2.066	118.679	16.310	1.00	28.80
	ATOM	953	N	LEU	120	-4.220	118.729	16.971	1.00	32.26
	ATOM	954	CA	LEU	120	-4.023	119.605	18.123	1.00	29.96
	ATOM	955	CB	LEU	120	-5.061	120.737	18.086	1.00	31.17
35	ATOM	956	CG	LEU	120	-5.022	121.998	18.959	1.00	30.26
	ATOM	957	CD1	LEU	120	-4.751	121.657	20.402	1.00	27.10
	ATOM	958	CD2	LEU	120	-3.971	122.926	18.420	1.00	30.96
	ATOM	959	C	LEU	120	-4.232	118.745	19.370	1.00	30.51
	ATOM	960	O	LEU	120	-5.339	118.271	19.633	1.00	32.88
40	ATOM	961	N	ALA	121	-3.168	118.549	20.140	1.00	27.40
	ATOM	962	CA	ALA	121	-3.241	117.728	21.341	1.00	25.55
	ATOM	963	CB	ALA	121	-2.338	116.516	21.177	1.00	19.41
	ATOM	964	C	ALA	121	-2.883	118.465	22.632	1.00	29.89
	ATOM	965	O	ALA	121	-1.943	119.256	22.671	1.00	35.88
45	ATOM	966	N	ILE	122	-3.650	118.199	23.685	1.00	30.33
	ATOM	967	CA	ILE	122	-3.406	118.797	24.990	1.00	30.27
	ATOM	968	CB	ILE	122	-4.692	119.357	25.605	1.00	29.27
	ATOM	969	CG2	ILE	122	-4.366	120.045	26.911	1.00	30.38
	ATOM	970	CG1	ILE	122	-5.347	120.344	24.635	1.00	26.03
50	ATOM	971	CD1	ILE	122	-6.653	120.942	25.125	1.00	29.21
	ATOM	972	C	ILE	122	-2.874	117.672	25.871	1.00	33.20
	ATOM	973	O	ILE	122	-3.587	116.712	26.155	1.00	33.30
	ATOM	974	N	PRO	123	-1.607	117.774	26.310	1.00	34.70
	ATOM	975	CD	PRO	123	-0.681	118.863	25.983	1.00	34.40
55	ATOM	976	CA	PRO	123	-0.937	116.778	27.157	1.00	36.49
	ATOM	977	CB	PRO	123	0.538	117.184	27.083	1.00	30.22
	ATOM	978	CG	PRO	123	0.614	118.125	25.915	1.00	36.72
	ATOM	979	C	PRO	123	-1.424	116.755	28.604	1.00	39.64
	ATOM	980	O	PRO	123	-0.623	116.919	29.530	1.00	40.48
60	ATOM	981	N	ARG	124	-2.724	116.551	28.798	1.00	44.58
	ATOM	982	CA	ARG	124	-3.301	116.510	30.134	1.00	48.04
	ATOM	983	CB	ARG	124	-3.715	117.908	30.576	1.00	52.91
	ATOM	984	CG	ARG	124	-2.595	118.930	30.504	1.00	60.50
	ATOM	985	CD	ARG	124	-2.167	119.290	31.910	1.00	72.57
65	ATOM	986	NE	ARG	124	-0.974	120.153	31.921	1.00	84.90
	ATOM	987	CZ	ARG	124	0.293	119.724	31.980	1.00	85.44

	ATOM	988	NH1	ARG	124	0.556	118.417	32.032	1.00	87.69
	ATOM	989	NH2	ARG	124	1.301	120.606	31.981	1.00	82.78
	ATOM	990	C	ARG	124	-4.496	115.573	30.204	1.00	48.39
	ATOM	991	O	ARG	124	-5.234	115.425	29.229	1.00	45.26
5	ATOM	992	N	GLU	125	-4.671	114.959	31.371	1.00	52.84
	ATOM	993	CA	GLU	125	-5.746	114.003	31.626	1.00	55.02
	ATOM	994	CB	GLU	125	-5.758	113.630	33.111	1.00	63.62
	ATOM	995	CG	GLU	125	-5.246	114.736	34.037	1.00	70.62
	ATOM	996	CD	GLU	125	-3.852	115.221	33.656	1.00	74.76
10	ATOM	997	OE1	GLU	125	-2.887	114.415	33.758	1.00	75.75
	ATOM	998	OE2	GLU	125	-3.724	116.404	33.254	1.00	80.71
	ATOM	999	C	GLU	125	-7.135	114.456	31.197	1.00	53.44
	ATOM	1000	O	GLU	125	-7.693	113.915	30.242	1.00	57.50
	ATOM	1001	N	ASN	126	-7.720	115.407	31.912	1.00	47.18
15	ATOM	1002	CA	ASN	126	-9.034	115.900	31.519	1.00	47.59
	ATOM	1003	CB	ASN	126	-10.127	115.425	32.473	1.00	51.03
	ATOM	1004	CG	ASN	126	-10.922	114.259	31.905	1.00	57.07
	ATOM	1005	OD1	ASN	126	-10.453	113.111	31.879	1.00	61.15
	ATOM	1006	ND2	ASN	126	-12.132	114.552	31.426	1.00	56.63
20	ATOM	1007	C	ASN	126	-8.972	117.402	31.501	1.00	47.06
	ATOM	1008	O	ASN	126	-9.558	118.086	32.345	1.00	45.61
	ATOM	1009	N	ALA	127	-8.234	117.904	30.520	1.00	43.50
	ATOM	1010	CA	ALA	127	-8.027	119.325	30.355	1.00	38.26
	ATOM	1011	CB	ALA	127	-7.390	119.588	29.011	1.00	37.97
25	ATOM	1012	C	ALA	127	-9.314	120.116	30.483	1.00	37.16
	ATOM	1013	O	ALA	127	-10.313	119.813	29.834	1.00	33.14
	ATOM	1014	N	GLN	128	-9.287	121.122	31.350	1.00	37.87
	ATOM	1015	CA	GLN	128	-10.436	121.998	31.522	1.00	37.19
	ATOM	1016	CB	GLN	128	-10.442	122.574	32.934	1.00	40.23
30	ATOM	1017	CG	GLN	128	-10.725	121.518	33.976	1.00	38.14
	ATOM	1018	CD	GLN	128	-12.006	120.779	33.670	1.00	39.51
	ATOM	1019	OE1	GLN	128	-13.101	121.347	33.743	1.00	40.36
	ATOM	1020	NE2	GLN	128	-11.881	119.505	33.299	1.00	41.53
	ATOM	1021	C	GLN	128	-10.252	123.084	30.465	1.00	35.55
35	ATOM	1022	O	GLN	128	-9.478	124.030	30.630	1.00	35.57
	ATOM	1023	N	ILE	129	-10.978	122.912	29.370	1.00	34.64
	ATOM	1024	CA	ILE	129	-10.888	123.779	28.203	1.00	29.60
	ATOM	1025	CB	ILE	129	-10.320	122.915	27.020	1.00	31.03
	ATOM	1026	CG2	ILE	129	-11.107	123.095	25.747	1.00	29.76
40	ATOM	1027	CG1	ILE	129	-8.850	123.220	26.837	1.00	26.37
	ATOM	1028	CD1	ILE	129	-8.045	122.897	28.039	1.00	34.70
	ATOM	1029	C	ILE	129	-12.225	124.406	27.821	1.00	29.55
	ATOM	1030	O	ILE	129	-13.278	123.868	28.129	1.00	33.42
	ATOM	1031	N	SER	130	-12.184	125.555	27.156	1.00	27.67
45	ATOM	1032	CA	SER	130	-13.411	126.198	26.698	1.00	25.00
	ATOM	1033	CB	SER	130	-13.273	127.710	26.694	1.00	22.09
	ATOM	1034	OG	SER	130	-14.381	128.297	26.027	1.00	20.89
	ATOM	1035	C	SER	130	-13.672	125.731	25.274	1.00	26.03
	ATOM	1036	O	SER	130	-12.757	125.684	24.472	1.00	30.77
50	ATOM	1037	N	LEU	131	-14.913	125.396	24.953	1.00	24.49
	ATOM	1038	CA	LEU	131	-15.230	124.925	23.614	1.00	28.07
	ATOM	1039	CB	LEU	131	-16.011	123.612	23.686	1.00	28.87
	ATOM	1040	CG	LEU	131	-15.183	122.328	23.730	1.00	30.76
	ATOM	1041	CD1	LEU	131	-14.079	122.430	24.740	1.00	28.98
55	ATOM	1042	CD2	LEU	131	-16.095	121.176	24.058	1.00	33.37
	ATOM	1043	C	LEU	131	-15.989	125.920	22.753	1.00	33.66
	ATOM	1044	O	LEU	131	-16.778	125.527	21.893	1.00	37.67
	ATOM	1045	N	ASP	132	-15.754	127.207	22.982	1.00	34.97
	ATOM	1046	CA	ASP	132	-16.401	128.248	22.190	1.00	34.31
60	ATOM	1047	CB	ASP	132	-16.456	129.566	22.962	1.00	42.68
	ATOM	1048	CG	ASP	132	-17.531	129.572	24.036	1.00	47.98
	ATOM	1049	OD1	ASP	132	-17.534	130.511	24.869	1.00	50.41
	ATOM	1050	OD2	ASP	132	-18.376	128.641	24.036	1.00	46.42
	ATOM	1051	C	ASP	132	-15.613	128.429	20.904	1.00	34.32
65	ATOM	1052	O	ASP	132	-14.391	128.521	20.922	1.00	32.87
	ATOM	1053	N	GLY	133	-16.329	128.478	19.789	1.00	34.53

	ATOM	1054	CA	GLY	133	-15.690	128.617	18.496	1.00	36.93
	ATOM	1055	C	GLY	133	-14.739	129.783	18.313	1.00	35.06
	ATOM	1056	O	GLY	133	-13.875	129.746	17.436	1.00	40.44
	ATOM	1057	N	ASP	134	-14.876	130.819	19.128	1.00	31.08
5	ATOM	1058	CA	ASP	134	-14.011	131.972	18.979	1.00	27.63
	ATOM	1059	CB	ASP	134	-14.801	133.258	19.236	1.00	28.43
	ATOM	1060	CG	ASP	134	-15.394	133.322	20.627	1.00	31.78
	ATOM	1061	OD1	ASP	134	-15.774	132.267	21.166	1.00	35.68
	ATOM	1062	OD2	ASP	134	-15.497	134.438	21.177	1.00	32.53
10	ATOM	1063	C	ASP	134	-12.753	131.922	19.827	1.00	25.08
	ATOM	1064	O	ASP	134	-11.786	132.581	19.518	1.00	19.71
	ATOM	1065	N	VAL	135	-12.739	131.115	20.875	1.00	26.84
	ATOM	1066	CA	VAL	135	-11.546	131.051	21.706	1.00	23.72
	ATOM	1067	CB	VAL	135	-11.902	131.077	23.194	1.00	25.13
15	ATOM	1068	CG1	VAL	135	-12.409	132.447	23.553	1.00	22.76
	ATOM	1069	CG2	VAL	135	-12.939	130.020	23.507	1.00	22.12
	ATOM	1070	C	VAL	135	-10.612	129.878	21.425	1.00	26.54
	ATOM	1071	O	VAL	135	-9.425	129.966	21.710	1.00	29.26
	ATOM	1072	N	THR	136	-11.128	128.777	20.882	1.00	26.39
20	ATOM	1073	CA	THR	136	-10.245	127.658	20.554	1.00	28.40
	ATOM	1074	CB	THR	136	-10.347	126.500	21.603	1.00	28.21
	ATOM	1075	OG1	THR	136	-11.127	125.427	21.079	1.00	37.13
	ATOM	1076	CG2	THR	136	-10.969	126.993	22.885	1.00	26.90
	ATOM	1077	C	THR	136	-10.512	127.160	19.126	1.00	31.60
25	ATOM	1078	O	THR	136	-11.608	126.696	18.799	1.00	27.29
	ATOM	1079	N	PHE	137	-9.493	127.291	18.276	1.00	31.45
	ATOM	1080	CA	PHE	137	-9.592	126.912	16.871	1.00	26.64
	ATOM	1081	CB	PHE	137	-10.012	128.133	16.067	1.00	26.84
	ATOM	1082	CG	PHE	137	-9.303	129.394	16.466	1.00	24.32
30	ATOM	1083	CD1	PHE	137	-8.067	129.721	15.916	1.00	27.77
	ATOM	1084	CD2	PHE	137	-9.871	130.262	17.384	1.00	23.70
	ATOM	1085	CE1	PHE	137	-7.414	130.889	16.271	1.00	27.32
	ATOM	1086	CE2	PHE	137	-9.221	131.432	17.745	1.00	26.31
	ATOM	1087	CZ	PHE	137	-7.991	131.745	17.185	1.00	29.43
35	ATOM	1088	C	PHE	137	-8.294	126.314	16.333	1.00	28.06
	ATOM	1089	O	PHE	137	-7.252	126.437	16.959	1.00	25.04
	ATOM	1090	N	PHE	138	-8.366	125.674	15.165	1.00	29.81
	ATOM	1091	CA	PHE	138	-7.210	125.003	14.578	1.00	29.93
	ATOM	1092	CB	PHE	138	-7.420	123.498	14.735	1.00	27.47
40	ATOM	1093	CG	PHE	138	-6.210	122.674	14.452	1.00	28.49
	ATOM	1094	CD1	PHE	138	-4.933	123.215	14.556	1.00	27.65
	ATOM	1095	CD2	PHE	138	-6.349	121.339	14.082	1.00	31.06
	ATOM	1096	CE1	PHE	138	-3.814	122.436	14.292	1.00	30.90
	ATOM	1097	CE2	PHE	138	-5.238	120.549	13.816	1.00	27.65
45	ATOM	1098	CZ	PHE	138	-3.968	121.097	13.919	1.00	27.41
	ATOM	1099	C	PHE	138	-6.928	125.400	13.116	1.00	30.91
	ATOM	1100	O	PHE	138	-7.799	125.340	12.256	1.00	24.94
	ATOM	1101	N	GLY	139	-5.667	125.772	12.879	1.00	36.94
	ATOM	1102	CA	GLY	139	-5.147	126.286	11.613	1.00	36.08
50	ATOM	1103	C	GLY	139	-5.072	125.640	10.252	1.00	36.48
	ATOM	1104	O	GLY	139	-6.057	125.113	9.769	1.00	43.61
	ATOM	1105	N	ALA	140	-3.905	125.763	9.616	1.00	36.60
	ATOM	1106	CA	ALA	140	-3.606	125.222	8.276	1.00	34.94
	ATOM	1107	CB	ALA	140	-4.170	123.820	8.136	1.00	36.94
55	ATOM	1108	C	ALA	140	-4.007	126.059	7.046	1.00	32.53
	ATOM	1109	O	ALA	140	-5.171	126.123	6.664	1.00	27.96
	ATOM	1110	N	LEU	141	-3.007	126.675	6.419	1.00	33.23
	ATOM	1111	CA	LEU	141	-3.180	127.503	5.218	1.00	36.11
	ATOM	1112	CB	LEU	141	-3.178	128.982	5.607	1.00	37.53
60	ATOM	1113	CG	LEU	141	-3.252	130.080	4.542	1.00	38.87
	ATOM	1114	CD1	LEU	141	-3.496	131.403	5.227	1.00	40.67
	ATOM	1115	CD2	LEU	141	-1.970	130.160	3.737	1.00	37.53
	ATOM	1116	C	LEU	141	-2.014	127.218	4.265	1.00	38.89
	ATOM	1117	O	LEU	141	-0.855	127.270	4.669	1.00	39.25
65	ATOM	1118	N	LYS	142	-2.303	126.927	3.001	1.00	42.86
	ATOM	1119	CA	LYS	142	-1.219	126.631	2.070	1.00	42.96

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	ATOM	1120	CB	LYS	142	-1.709	125.698	0.966	1.00	41.80
	ATOM	1121	CG	LYS	142	-0.626	125.387	-0.039	1.00	46.88
	ATOM	1122	CD	LYS	142	-0.916	124.136	-0.843	1.00	49.58
	ATOM	1123	CE	LYS	142	0.217	123.890	-1.826	1.00	51.07
5	ATOM	1124	NZ	LYS	142	0.041	122.632	-2.592	1.00	56.20
	ATOM	1125	C	LYS	142	-0.553	127.852	1.446	1.00	43.41
	ATOM	1126	O	LYS	142	-1.221	128.733	0.913	1.00	44.98
	ATOM	1127	N	LEU	143	0.775	127.893	1.513	1.00	42.81
	ATOM	1128	CA	LEU	143	1.542	129.002	0.950	1.00	40.65
10	ATOM	1129	CB	LEU	143	2.888	129.131	1.662	1.00	36.56
	ATOM	1130	CG	LEU	143	2.892	129.330	3.178	1.00	36.17
	ATOM	1131	CD1	LEU	143	4.327	129.369	3.670	1.00	37.95
	ATOM	1132	CD2	LEU	143	2.171	130.611	3.536	1.00	27.38
	ATOM	1133	C	LEU	143	1.802	128.772	-0.534	1.00	43.14
15	ATOM	1134	O	LEU	143	1.926	127.627	-0.981	1.00	46.28
	ATOM	1135	N	LEU	144	1.886	129.852	-1.300	1.00	42.63
	ATOM	1136	CA	LEU	144	2.150	129.727	-2.725	1.00	44.11
	ATOM	1137	CB	LEU	144	1.727	130.994	-3.463	1.00	44.88
	ATOM	1138	CG	LEU	144	0.225	131.270	-3.433	1.00	47.02
20	ATOM	1139	CD1	LEU	144	-0.069	132.587	-4.098	1.00	47.71
	ATOM	1140	CD2	LEU	144	-0.516	130.147	-4.135	1.00	47.87
	ATOM	1141	C	LEU	144	3.630	129.487	-2.955	1.00	45.66
	ATOM	1142	O	LEU	144	4.427	129.808	-2.055	1.00	46.37
	ATOM	1143	OXT	LEU	144	3.973	128.990	-4.044	1.00	50.66
25	END					60.588	44.144	42.385	0.00	0.00

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	ATOM	1	C	CYS	1	-5.491	39.770	-48.975 1.00101.54
5	ATOM	2	O	CYS	1	-6.060	39.762	-50.064 1.00102.94
	ATOM	3	CB	CYS	1	-5.049	41.915	-47.765 1.00 96.91
	ATOM	4	SG	CYS	1	-4.391	42.017	-46.070 1.00 92.39
	ATOM	5	N	CYS	1	-3.799	41.363	-49.827 1.00 93.44
	ATOM	6	CA	CYS	1	-4.430	40.797	-48.628 1.00 98.53
10	ATOM	7	N	SER	2	-5.724	38.920	-47.970 1.00104.82
	ATOM	8	CA	SER	2	-6.733	37.874	-47.875 1.00107.58
	ATOM	9	CB	SER	2	-6.818	36.973	-49.121 1.00109.37
	ATOM	10	OG	SER	2	-8.157	36.646	-49.431 1.00110.15
	ATOM	11	C	SER	2	-6.436	37.066	-46.585 1.00108.57
15	ATOM	12	O	SER	2	-6.569	37.627	-45.495 1.00109.82
	ATOM	13	N	GLN	3	-6.048	35.792	-46.667 1.00108.52
	ATOM	14	CA	GLN	3	-5.882	34.960	-45.460 1.00109.56
	ATOM	15	CB	GLN	3	-4.499	35.138	-44.804 1.00108.23
	ATOM	16	CG	GLN	3	-4.316	36.405	-43.997 1.00109.80
20	ATOM	17	CD	GLN	3	-3.469	37.365	-44.781 1.00110.35
	ATOM	18	OE1	GLN	3	-2.338	37.074	-45.156 1.00110.92
	ATOM	19	NE2	GLN	3	-3.831	38.586	-45.141 1.00110.59
	ATOM	20	C	GLN	3	-7.049	35.214	-44.473 1.00110.79
	ATOM	21	O	GLN	3	-6.847	35.559	-43.305 1.00114.50
25	ATOM	22	N	ASN	4	-8.277	35.028	-44.985 1.00109.87
	ATOM	23	CA	ASN	4	-9.600	35.143	-44.317 1.00107.78
	ATOM	24	CB	ASN	4	-9.872	33.893	-43.491 1.00109.08
	ATOM	25	CG	ASN	4	-8.736	33.511	-42.548 1.00109.99
	ATOM	26	OD1	ASN	4	-7.667	33.077	-42.981 1.00109.73
30	ATOM	27	ND2	ASN	4	-8.980	33.667	-41.245 1.00109.43
	ATOM	28	C	ASN	4	-9.793	36.424	-43.499 1.00105.15
	ATOM	29	O	ASN	4	-9.956	36.378	-42.290 1.00103.42
	ATOM	30	N	GLU	5	-9.750	37.597	-44.180 1.00101.11
	ATOM	31	CA	GLU	5	-9.920	38.925	-43.499 1.00 95.31
35	ATOM	32	CB	GLU	5	-8.603	39.374	-42.866 1.00 92.47
	ATOM	33	CG	GLU	5	-7.678	38.254	-42.418 1.00 86.82
	ATOM	34	CD	GLU	5	-6.266	38.536	-42.891 1.00 85.55
	ATOM	35	OE1	GLU	5	-6.112	38.955	-44.051 1.00 85.72
	ATOM	36	OE2	GLU	5	-5.337	38.339	-42.093 1.00 81.01
40	ATOM	37	C	GLU	5	-10.454	40.086	-44.375 1.00 92.66
	ATOM	38	O	GLU	5	-11.133	39.883	-45.366 1.00 94.43
	ATOM	39	N	TYR	6	-10.110	41.300	-43.945 1.00 85.19
	ATOM	40	CA	TYR	6	-10.526	42.537	-44.611 1.00 78.14
	ATOM	41	CB	TYR	6	-11.861	43.016	-44.072 1.00 76.59
45	ATOM	42	CG	TYR	6	-11.799	43.674	-42.729 1.00 73.94
	ATOM	43	CD1	TYR	6	-11.848	45.074	-42.649 1.00 73.49
	ATOM	44	CE1	TYR	6	-11.816	45.712	-41.426 1.00 73.13
	ATOM	45	CD2	TYR	6	-11.704	42.946	-41.540 1.00 72.45
	ATOM	46	CE2	TYR	6	-11.667	43.591	-40.294 1.00 72.38
50	ATOM	47	CZ	TYR	6	-11.730	44.963	-40.250 1.00 74.15
	ATOM	48	OH	TYR	6	-11.692	45.611	-39.038 1.00 73.32
	ATOM	49	C	TYR	6	-9.451	43.577	-44.394 1.00 75.43
	ATOM	50	O	TYR	6	-8.633	43.449	-43.464 1.00 76.93
	ATOM	51	N	PHE	7	-9.433	44.613	-45.229 1.00 69.11
55	ATOM	52	CA	PHE	7	-8.447	45.674	-45.063 1.00 62.54
	ATOM	53	CB	PHE	7	-7.848	46.035	-46.426 1.00 61.14
	ATOM	54	CG	PHE	7	-6.898	47.199	-46.340 1.00 61.11
	ATOM	55	CD1	PHE	7	-5.623	46.996	-45.827 1.00 59.90
	ATOM	56	CD2	PHE	7	-7.271	48.485	-46.728 1.00 63.23
60	ATOM	57	CE1	PHE	7	-4.729	48.061	-45.700 1.00 61.45
	ATOM	58	CE2	PHE	7	-6.388	49.566	-46.608 1.00 61.15
	ATOM	59	CZ	PHE	7	-5.117	49.355	-46.093 1.00 61.89
	ATOM	60	C	PHE	7	-9.065	46.908	-44.386 1.00 60.79
	ATOM	61	O	PHE	7	-9.673	47.744	-45.059 1.00 63.10
65	ATOM	62	N	ASP	8	-8.910	47.022	-43.063 1.00 57.06
	ATOM	63	CA	ASP	8	-9.456	48.160	-42.313 1.00 52.20

	ATOM	64	CB	ASP	8	-9.339	47.939	-40.808	1.00	46.52
	ATOM	65	CG	ASP	8	-10.114	48.937	-39.959	1.00	47.00
	ATOM	66	OD1	ASP	8	-10.206	50.111	-40.382	1.00	49.00
	ATOM	67	OD2	ASP	8	-10.609	48.544	-38.884	1.00	48.43
5	ATOM	68	C	ASP	8	-8.715	49.445	-42.710	1.00	53.02
	ATOM	69	O	ASP	8	-7.537	49.622	-42.383	1.00	53.25
	ATOM	70	N	SER	9	-9.409	50.334	-43.420	1.00	55.34
	ATOM	71	CA	SER	9	-8.824	51.608	-43.856	1.00	54.13
	ATOM	72	CB	SER	9	-9.769	52.344	-44.812	1.00	54.65
10	ATOM	73	OG	SER	9	-10.014	51.568	-45.971	1.00	54.09
	ATOM	74	C	SER	9	-8.503	52.531	-42.679	1.00	54.96
	ATOM	75	O	SER	9	-7.601	53.381	-42.772	1.00	54.11
	ATOM	76	N	LEU	10	-9.220	52.378	-41.570	1.00	49.64
	ATOM	77	CA	LEU	10	-8.930	53.212	-40.422	1.00	46.11
15	ATOM	78	CB	LEU	10	-9.936	52.994	-39.301	1.00	40.39
	ATOM	79	CG	LEU	10	-9.853	53.957	-38.138	1.00	34.58
	ATOM	80	CD1	LEU	10	-10.250	55.366	-38.583	1.00	33.04
	ATOM	81	CD2	LEU	10	-10.724	53.484	-36.972	1.00	33.82
	ATOM	82	C	LEU	10	-7.504	52.893	-39.971	1.00	49.82
20	ATOM	83	O	LEU	10	-6.718	53.788	-39.710	1.00	50.94
	ATOM	84	N	LEU	11	-7.181	51.603	-39.877	1.00	51.20
	ATOM	85	CA	LEU	11	-5.851	51.187	-39.435	1.00	51.39
	ATOM	86	CB	LEU	11	-5.926	49.824	-38.738	1.00	47.82
	ATOM	87	CG	LEU	11	-6.882	49.674	-37.546	1.00	46.28
25	ATOM	88	CD1	LEU	11	-6.772	48.272	-36.956	1.00	49.44
	ATOM	89	CD2	LEU	11	-6.598	50.719	-36.476	1.00	44.53
	ATOM	90	C	LEU	11	-4.793	51.127	-40.543	1.00	53.87
	ATOM	91	O	LEU	11	-3.602	51.199	-40.270	1.00	50.27
	ATOM	92	N	HIS	12	-5.232	50.980	-41.797	1.00	58.57
30	ATOM	93	CA	HIS	12	-4.339	50.864	-42.932	1.00	64.33
	ATOM	94	CB	HIS	12	-3.350	52.026	-42.998	1.00	65.68
	ATOM	95	CG	HIS	12	-4.032	53.388	-43.289	1.00	69.20
	ATOM	96	CD2	HIS	12	-5.148	53.691	-43.996	1.00	69.34
	ATOM	97	ND1	HIS	12	-3.561	54.584	-42.790	1.00	71.64
35	ATOM	98	CE1	HIS	12	-4.363	55.567	-43.171	1.00	72.29
	ATOM	99	NE2	HIS	12	-5.334	55.051	-43.904	1.00	70.37
	ATOM	100	C	HIS	12	-3.520	49.584	-42.828	1.00	66.69
	ATOM	101	O	HIS	12	-2.313	49.553	-43.090	1.00	67.11
	ATOM	102	N	ALA	13	-4.205	48.522	-42.414	1.00	69.57
40	ATOM	103	CA	ALA	13	-3.588	47.218	-42.218	1.00	71.62
	ATOM	104	CB	ALA	13	-2.980	47.103	-40.840	1.00	69.66
	ATOM	105	C	ALA	13	-4.648	46.141	-42.431	1.00	73.99
	ATOM	106	O	ALA	13	-5.793	46.453	-42.778	1.00	75.45
	ATOM	107	N	CYS	14	-4.283	44.880	-42.208	1.00	75.75
45	ATOM	108	CA	CYS	14	-5.229	43.790	-42.415	1.00	77.01
	ATOM	109	C	CYS	14	-5.681	43.097	-41.147	1.00	75.86
	ATOM	110	O	CYS	14	-4.889	42.521	-40.396	1.00	73.08
	ATOM	111	CB	CYS	14	-4.644	42.792	-43.417	1.00	80.96
	ATOM	112	SG	CYS	14	-5.227	43.010	-45.126	1.00	86.58
50	ATOM	113	N	ILE	15	-7.014	43.169	-40.940	1.00	75.89
	ATOM	114	CA	ILE	15	-7.624	42.593	-39.752	1.00	77.02
	ATOM	115	CB	ILE	15	-8.501	43.633	-39.016	1.00	76.68
	ATOM	116	CG2	ILE	15	-9.148	43.008	-37.784	1.00	76.37
	ATOM	117	CG1	ILE	15	-7.681	44.873	-38.617	1.00	76.28
55	ATOM	118	CD1	ILE	15	-6.600	44.607	-37.591	1.00	74.77
	ATOM	119	C	ILE	15	-8.462	41.326	-40.040	1.00	78.19
	ATOM	120	O	ILE	15	-9.251	41.254	-40.971	1.00	77.54
	ATOM	121	N	PRO	16	-8.235	40.335	-39.136	1.00	80.09
	ATOM	122	CD	PRO	16	-6.851	40.004	-38.725	1.00	79.64
60	ATOM	123	CA	PRO	16	-9.018	39.064	-39.202	1.00	82.16
	ATOM	124	CB	PRO	16	-8.442	38.269	-38.044	1.00	80.00
	ATOM	125	CG	PRO	16	-7.003	38.651	-38.092	1.00	80.17
	ATOM	126	C	PRO	16	-10.503	39.264	-39.289	1.00	84.28
	ATOM	127	O	PRO	16	-11.066	39.938	-38.429	1.00	84.12
65	ATOM	128	N	CYS	17	-11.174	38.711	-40.305	1.00	87.37
	ATOM	129	CA	CYS	17	-12.613	38.887	-40.419	1.00	88.33

	ATOM	130	CB	CYS	17	-13.173	37.993	-41.522	1.00	86.90
	ATOM	131	SG	CYS	17	-12.931	38.632	-43.206	1.00	84.56
	ATOM	132	C	CYS	17	-13.325	38.590	-39.094	1.00	89.64
	ATOM	133	O	CYS	17	-14.278	39.289	-38.712	1.00	89.77
5	ATOM	134	N	GLN	18	-12.855	37.548	-38.407	1.00	91.08
	ATOM	135	CA	GLN	18	-13.396	37.171	-37.134	1.00	91.16
	ATOM	136	CB	GLN	18	-12.417	36.321	-36.322	1.00	92.68
	ATOM	137	CG	GLN	18	-11.678	35.293	-37.170	1.00	98.31
	ATOM	138	CD	GLN	18	-10.353	34.807	-36.570	1.00	101.37
10	ATOM	139	OE1	GLN	18	-9.384	35.552	-36.500	1.00	102.74
	ATOM	140	NE2	GLN	18	-10.118	33.603	-36.083	1.00	103.76
	ATOM	141	C	GLN	18	-13.740	38.442	-36.353	1.00	89.18
	ATOM	142	O	GLN	18	-14.875	38.645	-35.918	1.00	89.64
	ATOM	143	N	LEU	19	-12.711	39.276	-36.190	1.00	87.56
15	ATOM	144	CA	LEU	19	-12.787	40.519	-35.443	1.00	85.33
	ATOM	145	CB	LEU	19	-11.527	41.311	-35.640	1.00	79.98
	ATOM	146	CG	LEU	19	-10.577	41.238	-34.464	1.00	77.91
	ATOM	147	CD1	LEU	19	-9.193	41.726	-34.860	1.00	78.13
	ATOM	148	CD2	LEU	19	-11.110	42.036	-33.286	1.00	72.50
20	ATOM	149	C	LEU	19	-13.989	41.391	-35.764	1.00	86.05
	ATOM	150	O	LEU	19	-14.644	41.907	-34.845	1.00	88.06
	ATOM	151	N	ARG	20	-14.300	41.559	-37.026	1.00	85.11
	ATOM	152	CA	ARG	20	-15.424	42.432	-37.387	1.00	83.33
	ATOM	153	CB	ARG	20	-15.246	42.957	-38.810	1.00	78.93
25	ATOM	154	CG	ARG	20	-14.773	44.405	-38.898	1.00	73.68
	ATOM	155	CD	ARG	20	-15.823	45.376	-38.389	1.00	67.23
	ATOM	156	NE	ARG	20	-16.323	46.246	-39.454	1.00	63.84
	ATOM	157	CZ	ARG	20	-15.541	46.931	-40.292	1.00	64.39
	ATOM	158	NH1	ARG	20	-14.220	46.836	-40.177	1.00	66.99
30	ATOM	159	NH2	ARG	20	-16.082	47.700	-41.223	1.00	60.63
	ATOM	160	C	ARG	20	-16.791	41.765	-37.228	1.00	84.21
	ATOM	161	O	ARG	20	-17.767	42.244	-37.805	1.00	86.28
	ATOM	162	N	CYS	21	-16.873	40.685	-36.464	1.00	85.02
	ATOM	163	CA	CYS	21	-18.175	40.056	-36.271	1.00	83.51
35	ATOM	164	CB	CYS	21	-18.081	38.566	-36.625	1.00	81.05
	ATOM	165	SG	CYS	21	-17.527	38.229	-38.327	1.00	73.08
	ATOM	166	C	CYS	21	-18.748	40.281	-34.849	1.00	84.26
	ATOM	167	O	CYS	21	-18.834	39.329	-34.085	1.00	84.48
	ATOM	168	N	SER	22	-19.132	41.551	-34.470	1.00	84.61
40	ATOM	169	CA	SER	22	-19.714	41.938	-33.147	1.00	85.07
	ATOM	170	C	SER	22	-21.168	42.355	-33.310	1.00	85.65
	ATOM	171	O	SER	22	-21.731	43.100	-32.512	1.00	86.77
	ATOM	172	CB	SER	22	-18.951	43.108	-32.511	1.00	20.00
	ATOM	173	OG	SER	22	-19.461	44.348	-32.963	1.00	20.00
45	ATOM	174	N	SER	23	-21.721	41.826	-34.368	1.00	84.19
	ATOM	175	CA	SER	23	-23.116	42.038	-34.681	1.00	82.44
	ATOM	176	C	SER	23	-23.351	43.088	-35.773	1.00	84.32
	ATOM	177	O	SER	23	-23.231	42.752	-36.952	1.00	81.24
	ATOM	178	CB	SER	23	-23.855	42.364	-33.383	1.00	78.00
50	ATOM	179	OG	SER	23	-24.881	43.315	-33.595	1.00	20.00
	ATOM	180	N	ASN	24	-23.671	44.307	-35.444	1.00	87.06
	ATOM	181	CA	ASN	24	-23.861	45.212	-36.560	1.00	88.17
	ATOM	182	C	ASN	24	-22.539	45.332	-37.333	1.00	88.45
	ATOM	183	O	ASN	24	-22.452	46.120	-38.258	1.00	84.89
55	ATOM	184	CB	ASN	24	-24.225	46.608	-36.106	0.00	86.84
	ATOM	185	CG	ASN	24	-25.400	46.626	-35.196	0.00	85.55
	ATOM	186	OD1	ASN	24	-26.250	45.743	-35.266	1.00	20.00
	ATOM	187	ND2	ASN	24	-25.478	47.628	-34.330	1.00	20.00
	ATOM	188	N	THR	25	-21.507	44.542	-36.935	1.00	89.49
60	ATOM	189	CA	THR	25	-20.179	44.757	-37.547	1.00	91.86
	ATOM	190	C	THR	25	-19.627	43.777	-38.638	1.00	95.75
	ATOM	191	O	THR	25	-18.405	43.627	-38.809	1.00	96.59
	ATOM	192	CB	THR	25	-19.188	44.954	-36.420	0.00	91.23
	ATOM	193	OG1	THR	25	-19.638	46.021	-35.584	1.00	20.00
65	ATOM	194	CG2	THR	25	-17.819	45.268	-36.983	1.00	20.00
	ATOM	195	N	PRO	26	-20.548	43.104	-39.365	1.00	98.41

	ATOM	196	CA	PRO	26	-20.212	42.246	-40.629	1.00	96.39
	ATOM	197	C	PRO	26	-19.784	42.914	-42.090	1.00	96.88
	ATOM	198	O	PRO	26	-20.712	43.211	-42.848	1.00	94.52
	ATOM	199	CB	PRO	26	-21.476	41.415	-40.733	0.00	95.29
5	ATOM	200	CG	PRO	26	-21.856	41.134	-39.310	0.00	92.34
	ATOM	201	CD	PRO	26	-21.265	42.211	-38.457	0.00	89.50
	ATOM	202	N	PRO	27	-18.441	43.180	-42.494	1.00	97.72
	ATOM	203	CA	PRO	27	-17.988	43.710	-43.904	1.00	98.94
	ATOM	204	C	PRO	27	-18.245	42.917	-45.257	1.00	100.71
10	ATOM	205	O	PRO	27	-18.567	41.734	-45.182	1.00	100.56
	ATOM	206	CB	PRO	27	-16.537	44.053	-43.649	0.00	97.33
	ATOM	207	CG	PRO	27	-16.575	44.598	-42.259	0.00	94.51
	ATOM	208	CD	PRO	27	-17.738	44.018	-41.540	0.00	92.58
	ATOM	209	N	LEU	28	-18.123	43.507	-46.512	1.00	15.00
15	ATOM	210	CA	LEU	28	-18.226	42.582	-47.688	1.00	15.00
	ATOM	211	C	LEU	28	-17.195	41.535	-47.272	1.00	15.00
	ATOM	212	O	LEU	28	-16.863	41.438	-46.090	1.00	15.00
	ATOM	213	CB	LEU	28	-17.865	43.242	-49.063	1.00	15.00
	ATOM	214	CG	LEU	28	-18.960	43.287	-50.174	1.00	15.00
20	ATOM	215	CD1	LEU	28	-18.564	42.401	-51.332	1.00	15.00
	ATOM	216	CD2	LEU	28	-20.319	42.875	-49.610	1.00	15.00
	ATOM	217	N	THR	29	-16.680	40.794	-48.239	1.00	15.00
	ATOM	218	CA	THR	29	-15.624	39.839	-47.927	1.00	15.00
	ATOM	219	C	THR	29	-15.657	39.468	-46.448	1.00	15.00
25	ATOM	220	O	THR	29	-14.659	39.045	-45.887	1.00	15.00
	ATOM	221	CB	THR	29	-14.261	40.421	-48.324	1.00	15.00
	ATOM	222	OG1	THR	29	-13.868	39.869	-49.577	1.00	20.00
	ATOM	223	CG2	THR	29	-13.227	40.091	-47.255	1.00	20.00
	ATOM	224	N	CYS	30	-16.850	39.635	-45.853	1.00	15.00
30	ATOM	225	CA	CYS	30	-16.980	39.354	-44.437	1.00	15.00
	ATOM	226	C	CYS	30	-18.303	38.656	-44.148	1.00	15.00
	ATOM	227	O	CYS	30	-18.388	37.876	-43.213	1.00	15.00
	ATOM	228	CB	CYS	30	-16.780	40.624	-43.622	1.00	15.00
	ATOM	229	SG	CYS	30	-15.060	41.215	-43.534	1.00	20.00
35	ATOM	230	N	GLN	31	-19.318	38.940	-44.964	1.00	15.00
	ATOM	231	CA	GLN	31	-20.589	38.281	-44.779	1.00	15.00
	ATOM	232	C	GLN	31	-20.448	36.760	-44.878	1.00	15.00
	ATOM	233	O	GLN	31	-20.919	36.009	-44.018	1.00	15.00
	ATOM	234	CB	GLN	31	-21.604	38.783	-45.774	1.00	15.00
40	ATOM	235	CG	GLN	31	-22.298	40.040	-45.265	1.00	20.00
	ATOM	236	CD	GLN	31	-23.400	40.517	-46.190	1.00	20.00
	ATOM	237	OE1	GLN	31	-23.896	39.760	-47.023	1.00	20.00
	ATOM	238	NE2	GLN	31	-23.935	41.733	-46.216	1.00	20.00
	ATOM	239	N	ARG	32	-19.798	36.317	-45.917	1.00	15.00
45	ATOM	240	CA	ARG	32	-19.534	34.911	-46.137	1.00	15.00
	ATOM	241	C	ARG	32	-18.766	34.288	-44.975	1.00	15.00
	ATOM	242	O	ARG	32	-19.046	33.153	-44.554	1.00	15.00
	ATOM	243	CB	ARG	32	-18.811	34.722	-47.488	1.00	15.00
	ATOM	244	CG	ARG	32	-19.607	35.185	-48.712	1.00	15.00
50	ATOM	245	CD	ARG	32	-18.838	34.881	-49.994	1.00	15.00
	ATOM	246	NE	ARG	32	-19.517	35.367	-51.189	1.00	20.00
	ATOM	247	CZ	ARG	32	-19.076	35.171	-52.426	1.00	20.00
	ATOM	248	NH1	ARG	32	-17.953	34.494	-52.629	1.00	20.00
	ATOM	249	NH2	ARG	32	-19.758	35.647	-53.459	1.00	20.00
55	ATOM	250	N	TYR	33	-17.814	35.066	-44.464	1.00	15.00
	ATOM	251	CA	TYR	33	-17.004	34.631	-43.340	1.00	15.00
	ATOM	252	C	TYR	33	-17.835	34.472	-42.036	1.00	15.00
	ATOM	253	O	TYR	33	-17.749	33.440	-41.384	1.00	15.00
	ATOM	254	CB	TYR	33	-15.885	35.607	-43.166	1.00	15.00
60	ATOM	255	CG	TYR	33	-15.043	35.761	-44.420	1.00	20.00
	ATOM	256	CD1	TYR	33	-15.321	36.723	-45.379	1.00	20.00
	ATOM	257	CD2	TYR	33	-13.941	34.939	-44.617	1.00	20.00
	ATOM	258	CE1	TYR	33	-14.526	36.866	-46.499	1.00	20.00
	ATOM	259	CE2	TYR	33	-13.138	35.075	-45.733	1.00	20.00
65	ATOM	260	CZ	TYR	33	-13.435	36.039	-46.671	1.00	20.00
	ATOM	261	OH	TYR	33	-12.639	36.179	-47.785	1.00	20.00

	ATOM	262	N	CYS	34	-18.634	35.489	-41.669	1.00	15.00
	ATOM	263	CA	CYS	34	-19.474	35.424	-40.482	1.00	15.00
	ATOM	264	C	CYS	34	-20.352	34.149	-40.545	1.00	15.00
	ATOM	265	O	CYS	34	-20.601	33.512	-39.521	1.00	15.00
5	ATOM	266	CB	CYS	34	-20.303	36.694	-40.334	1.00	15.00
	ATOM	267	SG	CYS	34	-19.340	38.199	-39.988	1.00	20.00
	ATOM	268	N	GLN	35	-20.795	33.804	-41.750	1.00	15.00
	ATOM	269	CA	GLN	35	-21.622	32.624	-42.080	1.00	15.00
	ATOM	270	CB	GLN	35	-22.092	32.776	-43.528	1.00	15.00
10	ATOM	271	CG	GLN	35	-23.270	31.928	-43.934	1.00	15.00
	ATOM	272	CD	GLN	35	-24.382	32.045	-42.934	1.00	15.00
	ATOM	273	OE1	GLN	35	-24.820	33.148	-42.616	1.00	15.00
	ATOM	274	NE2	GLN	35	-25.001	31.053	-42.325	1.00	15.00
	ATOM	275	C	GLN	35	-20.852	31.312	-41.960	1.00	15.00
15	ATOM	276	O	GLN	35	-21.340	30.346	-41.368	1.00	15.00
	ATOM	277	N	ALA	36	-19.647	31.324	-42.519	1.00	15.00
	ATOM	278	CA	ALA	36	-18.714	30.197	-42.472	1.00	15.00
	ATOM	279	C	ALA	36	-18.511	29.784	-41.003	1.00	15.00
	ATOM	280	O	ALA	36	-18.623	28.602	-40.668	1.00	15.00
20	ATOM	281	CB	ALA	36	-17.394	30.552	-43.110	1.00	15.00
	ATOM	282	N	SER	37	-18.187	30.790	-40.154	1.00	15.00
	ATOM	283	CA	SER	37	-17.929	30.468	-38.731	1.00	15.00
	ATOM	284	C	SER	37	-19.223	30.062	-38.066	1.00	15.00
	ATOM	285	O	SER	37	-19.249	29.134	-37.239	1.00	15.00
25	ATOM	286	CB	SER	37	-17.217	31.626	-38.013	1.00	15.00
	ATOM	287	OG	SER	37	-17.864	32.865	-38.245	1.00	20.00
	ATOM	288	N	VAL	38	-20.308	30.756	-38.422	1.00	15.00
	ATOM	289	CA	VAL	38	-21.606	30.436	-37.897	1.00	15.00
	ATOM	290	C	VAL	38	-21.958	28.960	-38.119	1.00	15.00
30	ATOM	291	O	VAL	38	-22.340	28.229	-37.186	1.00	15.00
	ATOM	292	CB	VAL	38	-22.697	31.319	-38.571	1.00	15.00
	ATOM	293	CG1	VAL	38	-24.101	30.802	-38.256	1.00	20.00
	ATOM	294	CG2	VAL	38	-22.565	32.767	-38.110	1.00	20.00
	ATOM	295	N	THR	39	-21.853	28.542	-39.392	1.00	15.00
35	ATOM	296	CA	THR	39	-22.178	27.169	-39.769	1.00	15.00
	ATOM	297	C	THR	39	-21.588	26.191	-38.763	1.00	15.00
	ATOM	298	O	THR	39	-21.832	24.983	-38.845	1.00	15.00
	ATOM	299	CB	THR	39	-21.621	26.871	-41.170	1.00	15.00
	ATOM	300	OG1	THR	39	-22.264	27.722	-42.134	1.00	20.00
40	ATOM	301	CG2	THR	39	-21.854	25.419	-41.515	1.00	20.00
	END					31.712	6.654	-112.989	0.00	0.00

TABLE 14

	111							
	ATOM	1	C	CYS	1	9.270	53.820	-3.718 1.00101.54
5	ATOM	2	O	CYS	1	10.492	53.933	-3.770 1.00102.94
	ATOM	3	CB	CYS	1	7.775	55.329	-5.041 1.00 96.91
	ATOM	4	SG	CYS	1	5.972	55.119	-5.179 1.00 92.39
	ATOM	5	N	CYS	1	9.037	56.206	-3.102 1.00 93.44
	ATOM	6	CA	CYS	1	8.352	55.026	-3.642 1.00 98.53
10	ATOM	7	N	SER	2	8.593	52.667	-3.733 1.00104.82
	ATOM	8	CA	SER	2	9.102	51.313	-3.906 1.00107.58
	ATOM	9	CB	SER	2	10.326	50.989	-3.029 1.00109.37
	ATOM	10	OG	SER	2	11.269	50.204	-3.728 1.00110.15
	ATOM	11	C	SER	2	7.920	50.343	-3.648 1.00108.57
15	ATOM	12	O	SER	2	6.973	50.345	-4.438 1.00109.82
	ATOM	13	N	GLN	3	7.942	49.527	-2.593 1.00108.52
	ATOM	14	CA	GLN	3	6.897	48.504	-2.391 1.00109.56
	ATOM	15	CB	GLN	3	5.647	49.066	-1.688 1.00108.23
	ATOM	16	CG	GLN	3	4.721	49.888	-2.558 1.00109.80
20	ATOM	17	CD	GLN	3	4.900	51.341	-2.225 1.00110.35
	ATOM	18	OE1	GLN	3	4.719	51.772	-1.091 1.00110.92
	ATOM	19	NE2	GLN	3	5.258	52.284	-3.082 1.00110.59
	ATOM	20	C	GLN	3	6.566	47.808	-3.735 1.00110.79
	ATOM	21	O	GLN	3	5.414	47.767	-4.177 1.00114.50
25	ATOM	22	N	ASN	4	7.618	47.261	-4.368 1.00109.87
	ATOM	23	CA	ASN	4	7.653	46.493	-5.640 1.00107.78
	ATOM	24	CB	ASN	4	7.195	45.063	-5.392 1.00109.08
	ATOM	25	CG	ASN	4	5.871	44.954	-4.643 1.00109.99
	ATOM	26	OD1	ASN	4	5.785	45.261	-3.453 1.00109.73
30	ATOM	27	ND2	ASN	4	4.833	44.502	-5.351 1.00109.43
	ATOM	28	C	ASN	4	6.896	47.146	-6.800 1.00105.15
	ATOM	29	O	ASN	4	5.923	46.602	-7.299 1.00103.42
	ATOM	30	N	GLU	5	7.345	48.355	-7.222 1.00101.11
	ATOM	31	CA	GLU	5	6.691	49.104	-8.347 1.00 95.31
35	ATOM	32	CB	GLU	5	5.464	49.863	-7.842 1.00 92.47
	ATOM	33	CG	GLU	5	4.751	49.235	-6.655 1.00 86.82
	ATOM	34	CD	GLU	5	4.460	50.297	-5.613 1.00 85.55
	ATOM	35	OE1	GLU	5	5.355	51.119	-5.354 1.00 85.72
	ATOM	36	OE2	GLU	5	3.342	50.291	-5.077 1.00 81.01
40	ATOM	37	C	GLU	5	7.587	50.102	-9.124 1.00 92.66
	ATOM	38	O	GLU	5	8.798	49.970	-9.178 1.00 94.43
	ATOM	39	N	TYR	6	6.920	51.091	-9.718 1.00 85.19
	ATOM	40	CA	TYR	6	7.568	52.130	-10.522 1.00 78.14
	ATOM	41	CB	TYR	6	7.683	51.695	-11.971 1.00 76.59
45	ATOM	42	CG	TYR	6	6.412	51.778	-12.757 1.00 73.94
	ATOM	43	CD1	TYR	6	6.218	52.857	-13.633 1.00 73.49
	ATOM	44	CE1	TYR	6	5.067	52.952	-14.390 1.00 73.13
	ATOM	45	CD2	TYR	6	5.406	50.813	-12.662 1.00 72.45
	ATOM	46	CE2	TYR	6	4.233	50.908	-13.426 1.00 72.38
50	ATOM	47	CZ	TYR	6	4.080	51.970	-14.284 1.00 74.15
	ATOM	48	OH	TYR	6	2.935	52.081	-15.038 1.00 73.32
	ATOM	49	C	TYR	6	6.756	53.399	-10.401 1.00 75.43
	ATOM	50	O	TYR	6	5.568	53.350	-10.031 1.00 76.93
	ATOM	51	N	PHE	7	7.366	54.540	-10.710 1.00 69.11
55	ATOM	52	CA	PHE	7	6.639	55.802	-10.649 1.00 62.54
	ATOM	53	CB	PHE	7	7.505	56.861	-9.959 1.00 61.14
	ATOM	54	CG	PHE	7	6.854	58.217	-9.958 1.00 61.11
	ATOM	55	CD1	PHE	7	5.819	58.472	-9.065 1.00 59.90
	ATOM	56	CD2	PHE	7	7.234	59.218	-10.852 1.00 63.23
60	ATOM	57	CE1	PHE	7	5.170	59.707	-9.063 1.00 61.45
	ATOM	58	CE2	PHE	7	6.594	60.463	-10.864 1.00 61.15
	ATOM	59	CZ	PHE	7	5.561	60.709	-9.971 1.00 61.89
	ATOM	60	C	PHE	7	6.212	56.267	-12.051 1.00 60.79
	ATOM	61	O	PHE	7	7.001	56.894	-12.763 1.00 63.10
65	ATOM	62	N	ASP	8	4.972	55.962	-12.446 1.00 57.06
	ATOM	63	CA	ASP	8	4.457	56.356	-13.763 1.00 52.20

	ATOM	64	CB	ASP	8	3.111	55.698	-14.052	1.00	46.52
	ATOM	65	CG	ASP	8	2.635	55.837	-15.490	1.00	47.00
	ATOM	66	OD1	ASP	8	2.924	56.891	-16.099	1.00	49.00
	ATOM	67	OD2	ASP	8	1.973	54.904	-15.990	1.00	48.43
5	ATOM	68	C	ASP	8	4.314	57.883	-13.826	1.00	53.02
	ATOM	69	O	ASP	8	3.449	58.466	-13.164	1.00	53.25
	ATOM	70	N	SER	9	5.170	58.527	-14.620	1.00	55.34
	ATOM	71	CA	SER	9	5.137	59.986	-14.779	1.00	54.13
	ATOM	72	CB	SER	9	6.344	60.474	-15.588	1.00	54.65
10	ATOM	73	OG	SER	9	7.554	60.145	-14.931	1.00	54.09
	ATOM	74	C	SER	9	3.859	60.464	-15.471	1.00	54.96
	ATOM	75	O	SER	9	3.420	61.608	-15.264	1.00	54.11
	ATOM	76	N	LEU	10	3.249	59.608	-16.286	1.00	49.64
	ATOM	77	CA	LEU	10	2.021	60.010	-16.940	1.00	46.11
15	ATOM	78	CB	LEU	10	1.546	58.961	-17.934	1.00	40.39
	ATOM	79	CG	LEU	10	0.388	59.365	-18.822	1.00	34.58
	ATOM	80	CD1	LEU	10	0.817	60.473	-19.786	1.00	33.04
	ATOM	81	CD2	LEU	10	-0.163	58.158	-19.585	1.00	33.82
	ATOM	82	C	LEU	10	0.982	60.264	-15.847	1.00	49.82
20	ATOM	83	O	LEU	10	0.284	61.264	-15.871	1.00	50.94
	ATOM	84	N	LEU	11	0.882	59.342	-14.890	1.00	51.20
	ATOM	85	CA	LEU	11	-0.094	59.476	-13.810	1.00	51.39
	ATOM	86	CB	LEU	11	-0.522	58.092	-13.307	1.00	47.82
	ATOM	87	CG	LEU	11	-1.091	57.097	-14.328	1.00	46.28
25	ATOM	88	CD1	LEU	11	-1.509	55.807	-13.630	1.00	49.44
	ATOM	89	CD2	LEU	11	-2.270	57.694	-15.083	1.00	44.53
	ATOM	90	C	LEU	11	0.375	60.319	-12.618	1.00	53.87
	ATOM	91	O	LEU	11	-0.439	60.842	-11.868	1.00	50.27
	ATOM	92	N	HIS	12	1.695	60.440	-12.438	1.00	58.57
30	ATOM	93	CA	HIS	12	2.271	61.170	-11.328	1.00	64.33
	ATOM	94	CB	HIS	12	1.734	62.598	-11.248	1.00	65.68
	ATOM	95	CG	HIS	12	2.169	63.480	-12.447	1.00	69.20
	ATOM	96	CD2	HIS	12	3.286	63.449	-13.214	1.00	69.34
	ATOM	97	ND1	HIS	12	1.383	64.490	-12.959	1.00	71.64
35	ATOM	98	CE1	HIS	12	1.993	65.040	-13.997	1.00	72.29
	ATOM	99	NE2	HIS	12	3.151	64.426	-14.173	1.00	70.37
	ATOM	100	C	HIS	12	1.926	60.486	-10.013	1.00	66.69
	ATOM	101	O	HIS	12	1.583	61.123	-9.011	1.00	67.11
	ATOM	102	N	ALA	13	2.003	59.159	-10.043	1.00	69.57
40	ATOM	103	CA	ALA	13	1.676	58.327	-8.893	1.00	71.62
	ATOM	104	CB	ALA	13	0.196	58.031	-8.837	1.00	69.66
	ATOM	105	C	ALA	13	2.482	57.034	-8.983	1.00	73.99
	ATOM	106	O	ALA	13	3.298	56.870	-9.898	1.00	75.45
	ATOM	107	N	CYS	14	2.247	56.109	-8.055	1.00	75.75
45	ATOM	108	CA	CYS	14	2.994	54.857	-8.054	1.00	77.01
	ATOM	109	C	CYS	14	2.176	53.634	-8.412	1.00	75.86
	ATOM	110	O	CYS	14	1.204	53.275	-7.741	1.00	73.08
	ATOM	111	CB	CYS	14	3.695	54.683	-6.705	1.00	80.96
	ATOM	112	SG	CYS	14	5.441	55.191	-6.693	1.00	86.58
50	ATOM	113	N	ILE	15	2.623	52.990	-9.512	1.00	75.89
	ATOM	114	CA	ILE	15	1.939	51.816	-10.028	1.00	77.02
	ATOM	115	CB	ILE	15	1.605	51.980	-11.530	1.00	76.68
	ATOM	116	CG2	ILE	15	0.904	50.733	-12.060	1.00	76.37
	ATOM	117	CG1	ILE	15	0.734	53.225	-11.775	1.00	76.28
55	ATOM	118	CD1	ILE	15	-0.649	53.155	-11.162	1.00	74.77
	ATOM	119	C	ILE	15	2.724	50.501	-9.818	1.00	78.19
	ATOM	120	O	ILE	15	3.920	50.401	-10.049	1.00	77.54
	ATOM	121	N	PRO	16	1.932	49.486	-9.377	1.00	80.09
	ATOM	122	CD	PRO	16	0.948	49.725	-8.296	1.00	79.64
60	ATOM	123	CA	PRO	16	2.497	48.115	-9.197	1.00	82.16
	ATOM	124	CB	PRO	16	1.296	47.332	-8.696	1.00	80.00
	ATOM	125	CG	PRO	16	0.611	48.336	-7.834	1.00	80.17
	ATOM	126	C	PRO	16	3.260	47.607	-10.386	1.00	84.28
	ATOM	127	O	PRO	16	2.706	47.580	-11.484	1.00	84.12
65	ATOM	128	N	CYS	17	4.525	47.206	-10.224	1.00	87.37
	ATOM	129	CA	CYS	17	5.292	46.710	-11.355	1.00	88.33

	ATOM	130	CB	CYS	17	6.616	46.116	-10.882	1.00	86.90
	ATOM	131	SG	CYS	17	7.903	47.345	-10.510	1.00	84.56
	ATOM	132	C	CYS	17	4.506	45.664	-12.155	1.00	89.64
	ATOM	133	O	CYS	17	4.553	45.643	-13.396	1.00	89.77
5	ATOM	134	N	GLN	18	3.793	44.800	-11.431	1.00	91.08
	ATOM	135	CA	GLN	18	2.980	43.788	-12.040	1.00	91.16
	ATOM	136	CB	GLN	18	1.895	43.275	-11.092	1.00	92.68
	ATOM	137	CG	GLN	18	2.392	43.095	-9.662	1.00	98.31
	ATOM	138	CD	GLN	18	1.287	43.113	-8.598	1.00	101.37
10	ATOM	139	OE1	GLN	18	0.685	44.147	-8.335	1.00	102.74
	ATOM	140	NE2	GLN	18	0.878	42.079	-7.887	1.00	103.76
	ATOM	141	C	GLN	18	2.328	44.374	-13.294	1.00	89.18
	ATOM	142	O	GLN	18	2.468	43.849	-14.401	1.00	89.64
	ATOM	143	N	LEU	19	1.606	45.474	-13.070	1.00	87.56
15	ATOM	144	CA	LEU	19	0.860	46.176	-14.098	1.00	85.33
	ATOM	145	CB	LEU	19	0.347	47.480	-13.557	1.00	79.98
	ATOM	146	CG	LEU	19	-1.125	47.451	-13.204	1.00	77.91
	ATOM	147	CD1	LEU	19	-1.491	48.637	-12.328	1.00	78.13
	ATOM	148	CD2	LEU	19	-1.983	47.425	-14.457	1.00	72.50
20	ATOM	149	C	LEU	19	1.620	46.428	-15.389	1.00	86.05
	ATOM	150	O	LEU	19	1.075	46.210	-16.483	1.00	88.06
	ATOM	151	N	ARG	20	2.852	46.865	-15.294	1.00	85.11
	ATOM	152	CA	ARG	20	3.610	47.169	-16.514	1.00	83.33
	ATOM	153	CB	ARG	20	4.711	48.182	-16.211	1.00	78.93
25	ATOM	154	CG	ARG	20	4.410	49.605	-16.672	1.00	73.68
	ATOM	155	CD	ARG	20	4.363	49.713	-18.185	1.00	67.23
	ATOM	156	NE	ARG	20	5.439	50.559	-18.705	1.00	63.84
	ATOM	157	CZ	ARG	20	5.724	51.778	-18.241	1.00	64.39
	ATOM	158	NH1	ARG	20	5.004	52.283	-17.244	1.00	66.99
30	ATOM	159	NH2	ARG	20	6.713	52.475	-18.777	1.00	60.63
	ATOM	160	C	ARG	20	4.194	45.930	-17.196	1.00	84.21
	ATOM	161	O	ARG	20	5.112	46.062	-18.004	1.00	86.28
	ATOM	162	N	CYS	21	3.680	44.748	-16.885	1.00	85.02
	ATOM	163	CA	CYS	21	4.200	43.559	-17.550	1.00	83.51
35	ATOM	164	CB	CYS	21	4.621	42.525	-16.496	1.00	81.05
	ATOM	165	SG	CYS	21	5.878	43.119	-15.320	1.00	73.08
	ATOM	166	C	CYS	21	3.208	42.964	-18.582	1.00	84.26
	ATOM	167	O	CYS	21	2.683	41.884	-18.349	1.00	84.48
	ATOM	168	N	SER	22	2.925	43.674	-19.732	1.00	84.61
40	ATOM	169	CA	SER	22	2.007	43.241	-20.831	1.00	85.07
	ATOM	170	C	SER	22	2.799	42.951	-22.098	1.00	85.65
	ATOM	171	O	SER	22	2.293	43.003	-23.215	1.00	86.77
	ATOM	172	CB	SER	22	0.965	44.320	-21.158	1.00	20.00
	ATOM	173	OG	SER	22	1.472	45.241	-22.104	1.00	20.00
45	ATOM	174	N	SER	23	4.042	42.639	-21.847	1.00	84.19
	ATOM	175	CA	SER	23	4.958	42.264	-22.900	1.00	82.44
	ATOM	176	C	SER	23	5.911	43.389	-23.319	1.00	84.32
	ATOM	177	O	SER	23	6.919	43.593	-22.640	1.00	81.24
	ATOM	178	CB	SER	23	4.143	41.718	-24.073	1.00	78.00
50	ATOM	179	OG	SER	23	4.716	42.077	-25.315	1.00	20.00
	ATOM	180	N	ASN	24	5.648	44.106	-24.374	1.00	87.06
	ATOM	181	CA	ASN	24	6.617	45.143	-24.667	1.00	88.17
	ATOM	182	C	ASN	24	6.648	46.137	-23.497	1.00	88.45
	ATOM	183	O	ASN	24	7.330	47.143	-23.580	1.00	84.89
55	ATOM	184	CB	ASN	24	6.246	45.937	-25.900	0.00	86.84
	ATOM	185	CG	ASN	24	6.010	45.075	-27.087	0.00	85.55
	ATOM	186	OD1	ASN	24	6.570	43.985	-27.181	1.00	20.00
	ATOM	187	ND2	ASN	24	5.185	45.538	-28.016	1.00	20.00
	ATOM	188	N	THR	25	5.892	45.845	-22.405	1.00	89.49
60	ATOM	189	CA	THR	25	5.771	46.862	-21.339	1.00	91.86
	ATOM	190	C	THR	25	6.563	46.718	-19.995	1.00	95.75
	ATOM	191	O	THR	25	6.146	47.234	-18.945	1.00	96.59
	ATOM	192	CB	THR	25	4.294	47.087	-21.096	0.00	91.23
	ATOM	193	OG1	THR	25	3.665	47.439	-22.330	1.00	20.00
65	ATOM	194	CG2	THR	25	4.099	48.186	-20.074	1.00	20.00
	ATOM	195	N	PRO	26	7.708	46.000	-20.045	1.00	98.41

	ATOM	196	CA	PRO	26	8.742	45.914	-18.874	1.00	96.39
	ATOM	197	C	PRO	26	9.742	47.175	-18.455	1.00	96.88
	ATOM	198	O	PRO	26	10.814	47.247	-19.062	1.00	94.52
	ATOM	199	CB	PRO	26	9.523	44.685	-19.295	0.00	95.29
5	ATOM	200	CG	PRO	26	8.492	43.773	-19.890	0.00	92.34
	ATOM	201	CD	PRO	26	7.352	44.618	-20.362	0.00	89.50
	ATOM	202	N	PRO	27	9.426	48.166	-17.478	1.00	97.72
	ATOM	203	CA	PRO	27	10.384	49.309	-16.977	1.00	98.94
	ATOM	204	C	PRO	27	11.771	49.029	-16.254	1.00	100.71
10	ATOM	205	O	PRO	27	11.985	47.895	-15.831	1.00	100.56
	ATOM	206	CB	PRO	27	9.434	50.178	-16.185	0.00	97.33
	ATOM	207	CG	PRO	27	8.181	50.107	-16.995	0.00	94.51
	ATOM	208	CD	PRO	27	8.170	48.834	-17.762	0.00	92.58
	ATOM	209	N	LEU	28	12.746	50.008	-16.086	1.00	15.00
15	ATOM	210	CA	LEU	28	13.920	49.631	-15.231	1.00	15.00
	ATOM	211	C	LEU	28	13.176	49.124	-13.998	1.00	15.00
	ATOM	212	O	LEU	28	11.997	48.782	-14.091	1.00	15.00
	ATOM	213	CB	LEU	28	14.877	50.823	-14.886	1.00	15.00
	ATOM	214	CG	LEU	28	16.364	50.738	-15.353	1.00	15.00
20	ATOM	215	CD1	LEU	28	17.280	50.621	-14.157	1.00	15.00
	ATOM	216	CD2	LEU	28	16.563	49.565	-16.311	1.00	15.00
	ATOM	217	N	THR	29	13.853	49.112	-12.862	1.00	15.00
	ATOM	218	CA	THR	29	13.178	48.728	-11.628	1.00	15.00
	ATOM	219	C	THR	29	11.943	47.887	-11.932	1.00	15.00
25	ATOM	220	O	THR	29	11.022	47.816	-11.134	1.00	15.00
	ATOM	221	CB	THR	29	12.813	49.981	-10.822	1.00	15.00
	ATOM	222	OG1	THR	29	13.778	50.166	-9.790	1.00	20.00
	ATOM	223	CG2	THR	29	11.422	49.822	-10.220	1.00	20.00
	ATOM	224	N	CYS	30	11.974	47.249	-13.114	1.00	15.00
30	ATOM	225	CA	CYS	30	10.830	46.457	-13.521	1.00	15.00
	ATOM	226	C	CYS	30	11.282	45.167	-14.194	1.00	15.00
	ATOM	227	O	CYS	30	10.590	44.165	-14.117	1.00	15.00
	ATOM	228	CB	CYS	30	9.889	47.287	-14.383	1.00	15.00
	ATOM	229	SG	CYS	30	8.930	48.543	-13.480	1.00	20.00
35	ATOM	230	N	GLN	31	12.448	45.208	-14.839	1.00	15.00
	ATOM	231	CA	GLN	31	12.963	44.011	-15.462	1.00	15.00
	ATOM	232	C	GLN	31	13.143	42.885	-14.441	1.00	15.00
	ATOM	233	O	GLN	31	12.697	41.751	-14.642	1.00	15.00
	ATOM	234	CB	GLN	31	14.262	44.292	-16.173	1.00	15.00
40	ATOM	235	CG	GLN	31	14.015	44.799	-17.588	1.00	20.00
	ATOM	236	CD	GLN	31	15.297	44.993	-18.373	1.00	20.00
	ATOM	237	OE1	GLN	31	16.341	44.445	-18.023	1.00	20.00
	ATOM	238	NE2	GLN	31	15.447	45.732	-19.468	1.00	20.00
	ATOM	239	N	ARG	32	13.787	43.204	-13.353	1.00	15.00
45	ATOM	240	CA	ARG	32	14.002	42.271	-12.266	1.00	15.00
	ATOM	241	C	ARG	32	12.688	41.717	-11.725	1.00	15.00
	ATOM	242	O	ARG	32	12.574	40.519	-11.414	1.00	15.00
	ATOM	243	CB	ARG	32	14.856	42.940	-11.168	1.00	15.00
	ATOM	244	CG	ARG	32	16.255	43.373	-11.618	1.00	15.00
50	ATOM	245	CD	ARG	32	17.039	43.946	-10.441	1.00	15.00
	ATOM	246	NE	ARG	32	18.354	44.444	-10.827	1.00	20.00
	ATOM	247	CZ	ARG	32	19.245	44.932	-9.972	1.00	20.00
	ATOM	248	NH1	ARG	32	18.958	44.987	-8.677	1.00	20.00
	ATOM	249	NH2	ARG	32	20.420	45.363	-10.408	1.00	20.00
55	ATOM	250	N	TYR	33	11.706	42.612	-11.641	1.00	15.00
	ATOM	251	CA	TYR	33	10.385	42.243	-11.163	1.00	15.00
	ATOM	252	C	TYR	33	9.660	41.260	-12.125	1.00	15.00
	ATOM	253	O	TYR	33	9.159	40.236	-11.680	1.00	15.00
	ATOM	254	CB	TYR	33	9.596	43.497	-10.958	1.00	15.00
60	ATOM	255	CG	TYR	33	10.272	44.463	-10.003	1.00	20.00
	ATOM	256	CD1	TYR	33	11.140	45.450	-10.447	1.00	20.00
	ATOM	257	CD2	TYR	33	10.006	44.389	-8.642	1.00	20.00
	ATOM	258	CE1	TYR	33	11.724	46.337	-9.565	1.00	20.00
	ATOM	259	CE2	TYR	33	10.583	45.274	-7.751	1.00	20.00
65	ATOM	260	CZ	TYR	33	11.441	46.246	-8.218	1.00	20.00
	ATOM	261	OH	TYR	33	12.019	47.130	-7.334	1.00	20.00

	ATOM	262	N	CYS	34	9.613	41.573	-13.431	1.00	15.00
	ATOM	263	CA	CYS	34	8.985	40.703	-14.414	1.00	15.00
	ATOM	264	C	CYS	34	9.592	39.283	-14.304	1.00	15.00
	ATOM	265	O	CYS	34	8.885	38.287	-14.462	1.00	15.00
5	ATOM	266	CB	CYS	34	9.116	41.286	-15.817	1.00	15.00
	ATOM	267	SG	CYS	34	8.195	42.831	-16.092	1.00	20.00
	ATOM	268	N	GLN	35	10.892	39.224	-14.029	1.00	15.00
	ATOM	269	CA	GLN	35	11.699	37.999	-13.847	1.00	15.00
	ATOM	270	CB	GLN	35	13.171	38.415	-13.799	1.00	15.00
10	ATOM	271	CG	GLN	35	14.176	37.320	-14.044	1.00	15.00
	ATOM	272	CD	GLN	35	13.822	36.534	-15.272	1.00	15.00
	ATOM	273	OE1	GLN	35	13.636	37.106	-16.345	1.00	15.00
	ATOM	274	NE2	GLN	35	13.690	35.226	-15.360	1.00	15.00
	ATOM	275	C	GLN	35	11.367	37.260	-12.554	1.00	15.00
15	ATOM	276	O	GLN	35	11.185	36.040	-12.553	1.00	15.00
	ATOM	277	N	ALA	36	11.278	38.037	-11.480	1.00	15.00
	ATOM	278	CA	ALA	36	10.912	37.549	-10.148	1.00	15.00
	ATOM	279	C	ALA	36	9.577	36.790	-10.250	1.00	15.00
	ATOM	280	O	ALA	36	9.463	35.664	-9.759	1.00	15.00
20	ATOM	281	CB	ALA	36	10.802	38.685	-9.162	1.00	15.00
	ATOM	282	N	SER	37	8.575	37.453	-10.878	1.00	15.00
	ATOM	283	CA	SER	37	7.244	36.809	-10.976	1.00	15.00
	ATOM	284	C	SER	37	7.324	35.634	-11.923	1.00	15.00
	ATOM	285	O	SER	37	6.713	34.580	-11.681	1.00	15.00
25	ATOM	286	CB	SER	37	6.156	37.824	-11.361	1.00	15.00
	ATOM	287	OG	SER	37	6.535	38.601	-12.482	1.00	20.00
	ATOM	288	N	VAL	38	8.079	35.810	-13.012	1.00	15.00
	ATOM	289	CA	VAL	38	8.273	34.753	-13.965	1.00	15.00
	ATOM	290	C	VAL	38	8.791	33.475	-13.295	1.00	15.00
30	ATOM	291	O	VAL	38	8.236	32.373	-13.466	1.00	15.00
	ATOM	292	CB	VAL	38	9.288	35.192	-15.062	1.00	15.00
	ATOM	293	CG1	VAL	38	9.738	34.001	-15.909	1.00	20.00
	ATOM	294	CG2	VAL	38	8.669	36.259	-15.959	1.00	20.00
	ATOM	295	N	THR	39	9.896	33.640	-12.548	1.00	15.00
35	ATOM	296	CA	THR	39	10.525	32.512	-11.865	1.00	15.00
	ATOM	297	C	THR	39	9.470	31.644	-11.197	1.00	15.00
	ATOM	298	O	THR	39	9.785	30.583	-10.649	1.00	15.00
	ATOM	299	CB	THR	39	11.514	33.032	-10.810	1.00	15.00
	ATOM	300	OG1	THR	39	12.572	33.759	-11.459	1.00	20.00
40	ATOM	301	CG2	THR	39	12.080	31.873	-10.024	1.00	20.00
	END					50.903	67.374	64.558	0.00	0.00

TABLE 15

	111								
	ATOM	1	C	CYS	1	-38.647	56.175	-15.856	1.00101.54
5	ATOM	2	O	CYS	1	-39.191	56.862	-14.995	1.00102.94
	ATOM	3	CB	CYS	1	-36.941	57.155	-17.403	1.00 96.91
	ATOM	4	SG	CYS	1	-35.944	56.182	-18.576	1.00 92.39
	ATOM	5	N	CYS	1	-39.327	57.773	-17.619	1.00 93.44
	ATOM	6	CA	CYS	1	-38.405	56.688	-17.263	1.00 98.53
10	ATOM	7	N	SER	2	-38.190	54.929	-15.684	1.00104.82
	ATOM	8	CA	SER	2	-38.137	54.139	-14.462	1.00107.58
	ATOM	9	CB	SER	2	-39.449	54.145	-13.656	1.00109.37
	ATOM	10	OG	SER	2	-39.202	54.204	-12.267	1.00110.15
	ATOM	11	C	SER	2	-37.700	52.706	-14.863	1.00108.57
15	ATOM	12	O	SER	2	-36.561	52.540	-15.305	1.00109.82
	ATOM	13	N	GLN	3	-38.546	51.684	-14.723	1.00108.52
	ATOM	14	CA	GLN	3	-38.120	50.293	-14.973	1.00109.56
	ATOM	15	CB	GLN	3	-38.202	49.908	-16.462	1.00108.23
	ATOM	16	CG	GLN	3	-37.089	50.441	-17.338	1.00109.80
20	ATOM	17	CD	GLN	3	-37.620	51.581	-18.160	1.00110.35
	ATOM	18	OE1	GLN	3	-38.570	51.443	-18.922	1.00110.92
	ATOM	19	NE2	GLN	3	-37.139	52.814	-18.157	1.00110.59
	ATOM	20	C	GLN	3	-36.714	50.050	-14.368	1.00110.79
	ATOM	21	O	GLN	3	-35.778	49.629	-15.054	1.00114.50
25	ATOM	22	N	ASN	4	-36.602	50.330	-13.059	1.00109.87
	ATOM	23	CA	ASN	4	-35.425	50.176	-12.164	1.00107.78
	ATOM	24	CB	ASN	4	-35.273	48.717	-11.757	1.00109.08
	ATOM	25	CG	ASN	4	-35.289	47.739	-12.926	1.00109.99
	ATOM	26	OD1	ASN	4	-36.322	47.525	-13.564	1.00109.73
30	ATOM	27	ND2	ASN	4	-34.131	47.134	-13.200	1.00109.43
	ATOM	28	C	ASN	4	-34.120	50.755	-12.720	1.00105.15
	ATOM	29	O	ASN	4	-33.166	50.033	-12.961	1.00103.42
	ATOM	30	N	GLU	5	-34.093	52.093	-12.948	1.00101.11
	ATOM	31	CA	GLU	5	-32.878	52.787	-13.495	1.00 95.31
35	ATOM	32	CB	GLU	5	-32.819	52.641	-15.016	1.00 92.47
	ATOM	33	CG	GLU	5	-33.452	51.377	-15.576	1.00 86.82
	ATOM	34	CD	GLU	5	-34.338	51.728	-16.754	1.00 85.55
	ATOM	35	OE1	GLU	5	-35.076	52.723	-16.653	1.00 85.72
	ATOM	36	OE2	GLU	5	-34.277	51.006	-17.760	1.00 81.01
40	ATOM	37	C	GLU	5	-32.729	54.290	-13.150	1.00 92.66
	ATOM	38	O	GLU	5	-33.241	54.775	-12.156	1.00 94.43
	ATOM	39	N	TYR	6	-31.998	54.984	-14.023	1.00 85.19
	ATOM	40	CA	TYR	6	-31.712	56.413	-13.877	1.00 78.14
	ATOM	41	CB	TYR	6	-30.454	56.629	-13.057	1.00 76.59
45	ATOM	42	CG	TYR	6	-29.173	56.374	-13.789	1.00 73.94
	ATOM	43	CD1	TYR	6	-28.430	57.462	-14.268	1.00 73.49
	ATOM	44	CE1	TYR	6	-27.233	57.263	-14.928	1.00 73.13
	ATOM	45	CD2	TYR	6	-28.678	55.085	-14.008	1.00 72.45
	ATOM	46	CE2	TYR	6	-27.464	54.878	-14.681	1.00 72.38
50	ATOM	47	CZ	TYR	6	-26.754	55.967	-15.126	1.00 74.15
	ATOM	48	OH	TYR	6	-25.564	55.782	-15.791	1.00 73.32
	ATOM	49	C	TYR	6	-31.568	57.011	-15.257	1.00 75.43
	ATOM	50	O	TYR	6	-31.323	56.280	-16.235	1.00 76.93
	ATOM	51	N	PHE	7	-31.708	58.330	-15.364	1.00 69.11
55	ATOM	52	CA	PHE	7	-31.549	58.983	-16.658	1.00 62.54
	ATOM	53	CB	PHE	7	-32.675	60.003	-16.858	1.00 61.14
	ATOM	54	CG	PHE	7	-32.511	60.790	-18.130	1.00 61.11
	ATOM	55	CD1	PHE	7	-32.829	60.191	-19.344	1.00 59.90
	ATOM	56	CD2	PHE	7	-32.016	62.093	-18.131	1.00 63.23
60	ATOM	57	CE1	PHE	7	-32.654	60.881	-20.544	1.00 61.45
	ATOM	58	CE2	PHE	7	-31.833	62.801	-19.326	1.00 61.15
	ATOM	59	CZ	PHE	7	-32.151	62.195	-20.534	1.00 61.89
	ATOM	60	C	PHE	7	-30.171	59.653	-16.778	1.00 60.79
	ATOM	61	O	PHE	7	-29.989	60.784	-16.320	1.00 63.10
65	ATOM	62	N	ASP	8	-29.206	58.960	-17.392	1.00 57.06
	ATOM	63	CA	ASP	8	-27.853	59.501	-17.566	1.00 52.20

	ATOM	64	CB	ASP	8	-26.893	58.436	-18.088	1.00	46.52
	ATOM	65	CG	ASP	8	-25.424	58.832	-18.045	1.00	47.00
	ATOM	66	OD1	ASP	8	-25.140	60.034	-18.239	1.00	49.00
	ATOM	67	OD2	ASP	8	-24.575	57.943	-17.828	1.00	48.43
5	ATOM	68	C	ASP	8	-27.891	60.689	-18.538	1.00	53.02
	ATOM	69	O	ASP	8	-28.121	60.516	-19.739	1.00	53.25
	ATOM	70	N	SER	9	-27.670	61.894	-18.011	1.00	55.34
	ATOM	71	CA	SER	9	-27.670	63.112	-18.830	1.00	54.13
	ATOM	72	CB	SER	9	-27.586	64.362	-17.948	1.00	54.65
10	ATOM	73	OG	SER	9	-28.699	64.436	-17.076	1.00	54.09
	ATOM	74	C	SER	9	-26.510	63.138	-19.828	1.00	54.96
	ATOM	75	O	SER	9	-26.604	63.781	-20.887	1.00	54.11
	ATOM	76	N	LEU	10	-25.419	62.448	-19.513	1.00	49.64
	ATOM	77	CA	LEU	10	-24.308	62.423	-20.441	1.00	46.11
15	ATOM	78	CB	LEU	10	-23.102	61.704	-19.856	1.00	40.39
	ATOM	79	CG	LEU	10	-21.820	61.797	-20.655	1.00	34.58
	ATOM	80	CD1	LEU	10	-21.298	63.235	-20.659	1.00	33.04
	ATOM	81	CD2	LEU	10	-20.764	60.832	-20.110	1.00	33.82
	ATOM	82	C	LEU	10	-24.797	61.751	-21.725	1.00	49.82
20	ATOM	83	O	LEU	10	-24.552	62.237	-22.816	1.00	50.94
	ATOM	84	N	LEU	11	-25.489	60.621	-21.582	1.00	51.20
	ATOM	85	CA	LEU	11	-25.986	59.886	-22.744	1.00	51.39
	ATOM	86	CB	LEU	11	-26.076	58.389	-22.426	1.00	47.82
	ATOM	87	CG	LEU	11	-24.808	57.678	-21.931	1.00	46.28
25	ATOM	88	CD1	LEU	11	-25.084	56.192	-21.725	1.00	49.44
	ATOM	89	CD2	LEU	11	-23.653	57.870	-22.904	1.00	44.53
	ATOM	90	C	LEU	11	-27.339	60.367	-23.281	1.00	53.87
	ATOM	91	O	LEU	11	-27.664	60.139	-24.439	1.00	50.27
	ATOM	92	N	HIS	12	-28.134	61.024	-22.430	1.00	58.57
30	ATOM	93	CA	HIS	12	-29.455	61.493	-22.795	1.00	64.33
	ATOM	94	CB	HIS	12	-29.422	62.364	-24.050	1.00	65.68
	ATOM	95	CG	HIS	12	-28.673	63.705	-23.840	1.00	69.20
	ATOM	96	CD2	HIS	12	-28.528	64.480	-22.737	1.00	69.34
	ATOM	97	ND1	HIS	12	-27.960	64.330	-24.841	1.00	71.64
35	ATOM	98	CE1	HIS	12	-27.399	65.430	-24.361	1.00	72.29
	ATOM	99	NE2	HIS	12	-27.729	65.543	-23.086	1.00	70.37
	ATOM	100	C	HIS	12	-30.369	60.313	-23.092	1.00	66.69
	ATOM	101	O	HIS	12	-31.149	60.311	-24.049	1.00	67.11
	ATOM	102	N	ALA	13	-30.239	59.289	-22.254	1.00	69.57
40	ATOM	103	CA	ALA	13	-31.001	58.056	-22.396	1.00	71.62
	ATOM	104	CB	ALA	13	-30.318	57.098	-23.343	1.00	69.66
	ATOM	105	C	ALA	13	-31.167	57.424	-21.016	1.00	73.99
	ATOM	106	O	ALA	13	-30.736	58.001	-20.010	1.00	75.45
	ATOM	107	N	CYS	14	-31.770	56.238	-20.963	1.00	75.75
45	ATOM	108	CA	CYS	14	-31.992	55.580	-19.681	1.00	77.01
	ATOM	109	C	CYS	14	-31.162	54.333	-19.459	1.00	75.86
	ATOM	110	O	CYS	14	-31.250	53.347	-20.196	1.00	73.08
	ATOM	111	CB	CYS	14	-33.485	55.294	-19.507	1.00	80.96
	ATOM	112	SG	CYS	14	-34.377	56.525	-18.509	1.00	86.58
50	ATOM	113	N	ILE	15	-30.344	54.414	-18.387	1.00	75.89
	ATOM	114	CA	ILE	15	-29.444	53.325	-18.043	1.00	77.02
	ATOM	115	CB	ILE	15	-27.990	53.831	-17.887	1.00	76.68
	ATOM	116	CG2	ILE	15	-27.064	52.681	-17.507	1.00	76.37
	ATOM	117	CG1	ILE	15	-27.496	54.511	-19.176	1.00	76.28
55	ATOM	118	CD1	ILE	15	-27.369	53.585	-20.367	1.00	74.77
	ATOM	119	C	ILE	15	-29.861	52.560	-16.766	1.00	78.19
	ATOM	120	O	ILE	15	-30.215	53.127	-15.742	1.00	77.54
	ATOM	121	N	PRO	16	-29.764	51.210	-16.912	1.00	80.09
	ATOM	122	CD	PRO	16	-30.268	50.556	-18.141	1.00	79.64
60	ATOM	123	CA	PRO	16	-30.044	50.307	-15.755	1.00	82.16
	ATOM	124	CB	PRO	16	-29.830	48.931	-16.360	1.00	80.00
	ATOM	125	CG	PRO	16	-30.365	49.112	-17.738	1.00	80.17
	ATOM	126	C	PRO	16	-29.312	50.678	-14.498	1.00	84.28
	ATOM	127	O	PRO	16	-28.087	50.783	-14.529	1.00	84.12
65	ATOM	128	N	CYS	17	-30.011	50.894	-13.379	1.00	87.37
	ATOM	129	CA	CYS	17	-29.332	51.257	-12.145	1.00	88.33

	ATOM	130	CB	CYS	17	-30.310	51.236	-10.974	1.00	86.90
	ATOM	131	SG	CYS	17	-31.376	52.704	-10.854	1.00	84.56
	ATOM	132	C	CYS	17	-28.150	50.324	-11.856	1.00	89.64
	ATOM	133	O	CYS	17	-27.085	50.769	-11.396	1.00	89.77
5	ATOM	134	N	GLN	18	-28.354	49.034	-12.124	1.00	91.08
	ATOM	135	CA	GLN	18	-27.330	48.048	-11.939	1.00	91.16
	ATOM	136	CB	GLN	18	-27.590	46.786	-12.763	1.00	92.68
	ATOM	137	CG	GLN	18	-29.056	46.369	-12.764	1.00	98.31
	ATOM	138	CD	GLN	18	-29.465	45.486	-13.949	1.00	101.37
10	ATOM	139	OE1	GLN	18	-29.519	45.943	-15.084	1.00	102.74
	ATOM	140	NE2	GLN	18	-29.783	44.206	-13.889	1.00	103.76
	ATOM	141	C	GLN	18	-25.986	48.657	-12.347	1.00	89.18
	ATOM	142	O	GLN	18	-25.029	48.690	-11.570	1.00	89.64
	ATOM	143	N	LEU	19	-25.957	49.124	-13.596	1.00	87.56
15	ATOM	144	CA	LEU	19	-24.778	49.702	-14.217	1.00	85.33
	ATOM	145	CB	LEU	19	-25.146	50.321	-15.535	1.00	79.98
	ATOM	146	CG	LEU	19	-24.755	49.477	-16.729	1.00	77.91
	ATOM	147	CD1	LEU	19	-25.473	49.952	-17.982	1.00	78.13
	ATOM	148	CD2	LEU	19	-23.250	49.494	-16.935	1.00	72.50
20	ATOM	149	C	LEU	19	-24.036	50.721	-13.369	1.00	86.05
	ATOM	150	O	LEU	19	-22.799	50.674	-13.284	1.00	88.06
	ATOM	151	N	ARG	20	-24.749	51.623	-12.739	1.00	85.11
	ATOM	152	CA	ARG	20	-24.073	52.658	-11.946	1.00	83.33
	ATOM	153	CB	ARG	20	-24.968	53.888	-11.817	1.00	78.93
25	ATOM	154	CG	ARG	20	-24.573	55.058	-12.714	1.00	73.68
	ATOM	155	CD	ARG	20	-23.239	55.658	-12.308	1.00	67.23
	ATOM	156	NE	ARG	20	-23.384	57.033	-11.827	1.00	63.84
	ATOM	157	CZ	ARG	20	-24.053	57.988	-12.477	1.00	64.39
	ATOM	158	NH1	ARG	20	-24.637	57.703	-13.637	1.00	66.99
30	ATOM	159	NH2	ARG	20	-24.126	59.208	-11.969	1.00	60.63
	ATOM	160	C	ARG	20	-23.623	52.174	-10.567	1.00	84.21
	ATOM	161	O	ARG	20	-23.365	53.000	-9.692	1.00	86.28
	ATOM	162	N	CYS	21	-23.527	50.868	-10.365	1.00	85.02
	ATOM	163	CA	CYS	21	-23.065	50.388	-9.067	1.00	83.51
35	ATOM	164	CB	CYS	21	-24.077	49.380	-8.504	1.00	81.05
	ATOM	165	SG	CYS	21	-25.764	50.037	-8.306	1.00	73.08
	ATOM	166	C	CYS	21	-21.630	49.804	-9.116	1.00	84.26
	ATOM	167	O	CYS	21	-21.471	48.602	-8.957	1.00	84.48
	ATOM	168	N	SER	22	-20.566	50.651	-9.356	1.00	84.61
40	ATOM	169	CA	SER	22	-19.125	50.257	-9.422	1.00	85.07
	ATOM	170	C	SER	22	-18.362	50.844	-8.243	1.00	85.65
	ATOM	171	O	SER	22	-17.150	51.042	-8.279	1.00	86.77
	ATOM	172	CB	SER	22	-18.459	50.750	-10.714	1.00	20.00
	ATOM	173	OG	SER	22	-17.969	52.068	-10.557	1.00	20.00
45	ATOM	174	N	SER	23	-19.137	51.091	-7.222	1.00	84.19
	ATOM	175	CA	SER	23	-18.610	51.594	-5.974	1.00	82.44
	ATOM	176	C	SER	23	-18.815	53.100	-5.779	1.00	84.32
	ATOM	177	O	SER	23	-19.907	53.501	-5.375	1.00	81.24
	ATOM	178	CB	SER	23	-17.141	51.182	-5.873	1.00	78.00
50	ATOM	179	OG	SER	23	-16.365	52.182	-5.242	1.00	20.00
	ATOM	180	N	ASN	24	-17.843	53.928	-6.040	1.00	87.06
	ATOM	181	CA	ASN	24	-18.156	55.326	-5.825	1.00	88.17
	ATOM	182	C	ASN	24	-19.299	55.730	-6.768	1.00	88.45
	ATOM	183	O	ASN	24	-19.656	56.894	-6.818	1.00	84.89
55	ATOM	184	CB	ASN	24	-16.986	56.228	-6.152	0.00	86.84
	ATOM	185	CG	ASN	24	-15.746	55.840	-5.432	0.00	85.55
	ATOM	186	OD1	ASN	24	-15.813	55.259	-4.352	1.00	20.00
	ATOM	187	ND2	ASN	24	-14.591	56.153	-6.005	1.00	20.00
	ATOM	188	N	THR	25	-19.864	54.751	-7.523	1.00	89.49
60	ATOM	189	CA	THR	25	-20.846	55.137	-8.558	1.00	91.86
	ATOM	190	C	THR	25	-22.381	54.920	-8.333	1.00	95.75
	ATOM	191	O	THR	25	-23.157	54.768	-9.292	1.00	96.59
	ATOM	192	CB	THR	25	-20.383	54.535	-9.867	0.00	91.23
	ATOM	193	OG1	THR	25	-19.044	54.959	-10.129	1.00	20.00
65	ATOM	194	CG2	THR	25	-21.301	54.968	-10.990	1.00	20.00
	ATOM	195	N	PRO	26	-22.805	54.899	-7.048	1.00	98.41

	ATOM	196	CA	PRO	26	-24.309	54.904	-6.619	1.00	96.39
	ATOM	197	C	PRO	26	-25.283	56.246	-6.748	1.00	96.88
	ATOM	198	O	PRO	26	-25.267	57.026	-5.791	1.00	94.52
	ATOM	199	CB	PRO	26	-24.181	54.429	-5.185	0.00	95.29
5	ATOM	200	CG	PRO	26	-23.076	53.417	-5.225	0.00	92.34
	ATOM	201	CD	PRO	26	-22.212	53.727	-6.405	0.00	89.50
	ATOM	202	N	PRO	27	-26.092	56.551	-7.882	1.00	97.72
	ATOM	203	CA	PRO	27	-27.105	57.749	-8.000	1.00	98.94
	ATOM	204	C	PRO	27	-28.364	57.923	-7.047	1.00	100.71
10	ATOM	205	O	PRO	27	-28.715	56.959	-6.371	1.00	100.56
	ATOM	206	CB	PRO	27	-27.439	57.722	-9.475	0.00	97.33
	ATOM	207	CG	PRO	27	-26.130	57.359	-10.095	0.00	94.51
	ATOM	208	CD	PRO	27	-25.319	56.598	-9.110	0.00	92.58
	ATOM	209	N	LEU	28	-29.076	59.115	-6.947	1.00	15.00
15	ATOM	210	CA	LEU	28	-30.340	59.062	-6.140	1.00	15.00
	ATOM	211	C	LEU	28	-31.013	57.865	-6.808	1.00	15.00
	ATOM	212	O	LEU	28	-30.336	57.065	-7.455	1.00	15.00
	ATOM	213	CB	LEU	28	-31.221	60.354	-6.235	1.00	15.00
	ATOM	214	CG	LEU	28	-31.509	61.154	-4.926	1.00	15.00
20	ATOM	215	CD1	LEU	28	-32.976	61.069	-4.575	1.00	15.00
	ATOM	216	CD2	LEU	28	-30.641	60.640	-3.778	1.00	15.00
	ATOM	217	N	THR	29	-32.325	57.774	-6.676	1.00	15.00
	ATOM	218	CA	THR	29	-33.043	56.708	-7.365	1.00	15.00
	ATOM	219	C	THR	29	-32.103	55.554	-7.694	1.00	15.00
25	ATOM	220	O	THR	29	-32.358	54.779	-8.602	1.00	15.00
	ATOM	221	CB	THR	29	-33.708	57.262	-8.632	1.00	15.00
	ATOM	222	OG1	THR	29	-35.087	57.503	-8.365	1.00	20.00
	ATOM	223	CG2	THR	29	-33.558	56.264	-9.773	1.00	20.00
	ATOM	224	N	CYS	30	-31.017	55.471	-6.906	1.00	15.00
30	ATOM	225	CA	CYS	30	-30.035	54.435	-7.162	1.00	15.00
	ATOM	226	C	CYS	30	-29.525	53.846	-5.854	1.00	15.00
	ATOM	227	O	CYS	30	-29.158	52.683	-5.812	1.00	15.00
	ATOM	228	CB	CYS	30	-28.924	54.965	-8.057	1.00	15.00
	ATOM	229	SG	CYS	30	-29.392	55.207	-9.799	1.00	20.00
35	ATOM	230	N	GLN	31	-29.518	54.658	-4.796	1.00	15.00
	ATOM	231	CA	GLN	31	-29.091	54.155	-3.512	1.00	15.00
	ATOM	232	C	GLN	31	-29.949	52.970	-3.063	1.00	15.00
	ATOM	233	O	GLN	31	-29.442	51.915	-2.669	1.00	15.00
	ATOM	234	CB	GLN	31	-29.114	55.247	-2.473	1.00	15.00
40	ATOM	235	CG	GLN	31	-27.814	56.040	-2.480	1.00	20.00
	ATOM	236	CD	GLN	31	-27.754	57.080	-1.379	1.00	20.00
	ATOM	237	OE1	GLN	31	-28.498	57.007	-0.401	1.00	20.00
	ATOM	238	NE2	GLN	31	-26.947	58.134	-1.333	1.00	20.00
	ATOM	239	N	ARG	32	-31.239	53.147	-3.132	1.00	15.00
45	ATOM	240	CA	ARG	32	-32.192	52.111	-2.791	1.00	15.00
	ATOM	241	C	ARG	32	-31.985	50.852	-3.627	1.00	15.00
	ATOM	242	O	ARG	32	-32.076	49.721	-3.118	1.00	15.00
	ATOM	243	CB	ARG	32	-33.628	52.666	-2.910	1.00	15.00
	ATOM	244	CG	ARG	32	-33.943	53.836	-1.972	1.00	15.00
50	ATOM	245	CD	ARG	32	-35.404	54.252	-2.113	1.00	15.00
	ATOM	246	NE	ARG	32	-35.742	55.412	-1.297	1.00	20.00
	ATOM	247	CZ	ARG	32	-36.963	55.925	-1.203	1.00	20.00
	ATOM	248	NH1	ARG	32	-37.965	55.375	-1.878	1.00	20.00
	ATOM	249	NH2	ARG	32	-37.184	56.982	-0.434	1.00	20.00
55	ATOM	250	N	TYR	33	-31.688	51.081	-4.903	1.00	15.00
	ATOM	251	CA	TYR	33	-31.440	49.990	-5.830	1.00	15.00
	ATOM	252	C	TYR	33	-30.152	49.193	-5.478	1.00	15.00
	ATOM	253	O	TYR	33	-30.195	47.972	-5.403	1.00	15.00
	ATOM	254	CB	TYR	33	-31.378	50.556	-7.212	1.00	15.00
60	ATOM	255	CG	TYR	33	-32.636	51.318	-7.588	1.00	20.00
	ATOM	256	CD1	TYR	33	-32.764	52.681	-7.369	1.00	20.00
	ATOM	257	CD2	TYR	33	-33.692	50.651	-8.196	1.00	20.00
	ATOM	258	CE1	TYR	33	-33.905	53.362	-7.744	1.00	20.00
	ATOM	259	CE2	TYR	33	-34.838	51.323	-8.577	1.00	20.00
65	ATOM	260	CZ	TYR	33	-34.939	52.678	-8.348	1.00	20.00
	ATOM	261	OH	TYR	33	-36.079	53.352	-8.726	1.00	20.00

	ATOM	262	N	CYS	34	-29.021	49.886	-5.261	1.00	15.00
	ATOM	263	CA	CYS	34	-27.772	49.235	-4.896	1.00	15.00
	ATOM	264	C	CYS	34	-28.005	48.336	-3.656	1.00	15.00
	ATOM	265	O	CYS	34	-27.427	47.254	-3.551	1.00	15.00
5	ATOM	266	CB	CYS	34	-26.669	50.263	-4.670	1.00	15.00
	ATOM	267	SG	CYS	34	-26.156	51.172	-6.161	1.00	20.00
	ATOM	268	N	GLN	35	-28.856	48.806	-2.747	1.00	15.00
	ATOM	269	CA	GLN	35	-29.267	48.134	-1.497	1.00	15.00
	ATOM	270	CB	GLN	35	-30.050	49.148	-0.661	1.00	15.00
10	ATOM	271	CG	GLN	35	-30.196	48.825	0.804	1.00	15.00
	ATOM	272	CD	GLN	35	-28.872	48.458	1.405	1.00	15.00
	ATOM	273	OE1	GLN	35	-27.907	49.211	1.292	1.00	15.00
	ATOM	274	NE2	GLN	35	-28.594	47.371	2.096	1.00	15.00
	ATOM	275	C	GLN	35	-30.162	46.923	-1.745	1.00	15.00
15	ATOM	276	O	GLN	35	-29.948	45.851	-1.171	1.00	15.00
	ATOM	277	N	ALA	36	-31.142	47.128	-2.618	1.00	15.00
	ATOM	278	CA	ALA	36	-32.081	46.090	-3.050	1.00	15.00
	ATOM	279	C	ALA	36	-31.279	44.882	-3.567	1.00	15.00
	ATOM	280	O	ALA	36	-31.536	43.746	-3.162	1.00	15.00
20	ATOM	281	CB	ALA	36	-33.011	46.606	-4.120	1.00	15.00
	ATOM	282	N	SER	37	-30.326	45.166	-4.488	1.00	15.00
	ATOM	283	CA	SER	37	-29.541	44.053	-5.071	1.00	15.00
	ATOM	284	C	SER	37	-28.627	43.477	-4.015	1.00	15.00
	ATOM	285	O	SER	37	-28.437	42.251	-3.937	1.00	15.00
25	ATOM	286	CB	SER	37	-28.796	44.494	-6.341	1.00	15.00
	ATOM	287	OG	SER	37	-28.078	45.697	-6.138	1.00	20.00
	ATOM	288	N	VAL	38	-28.050	44.361	-3.194	1.00	15.00
	ATOM	289	CA	VAL	38	-27.198	43.937	-2.119	1.00	15.00
	ATOM	290	C	VAL	38	-27.893	42.912	-1.215	1.00	15.00
30	ATOM	291	O	VAL	38	-27.365	41.821	-0.931	1.00	15.00
	ATOM	292	CB	VAL	38	-26.766	45.158	-1.255	1.00	15.00
	ATOM	293	CG1	VAL	38	-26.112	44.709	0.052	1.00	20.00
	ATOM	294	CG2	VAL	38	-25.801	46.044	-2.036	1.00	20.00
	ATOM	295	N	THR	39	-29.087	43.303	-0.738	1.00	15.00
35	ATOM	296	CA	THR	39	-29.863	42.449	0.158	1.00	15.00
	ATOM	297	C	THR	39	-29.856	41.013	-0.344	1.00	15.00
	ATOM	298	O	THR	39	-30.372	40.112	0.324	1.00	15.00
	ATOM	299	CB	THR	39	-31.309	42.963	0.240	1.00	15.00
	ATOM	300	OG1	THR	39	-31.320	44.278	0.820	1.00	20.00
40	ATOM	301	CG2	THR	39	-32.142	42.016	1.072	1.00	20.00
	END					-119.496	62.638	-15.481	0.00	0.00

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	ATOM	1	C	CYS	1	56.065	82.669	3.315 1.00101.54
5	ATOM	2	O	CYS	1	56.592	82.165	2.326 1.00102.94
	ATOM	3	CB	CYS	1	54.032	84.113	3.099 1.00 96.91
	ATOM	4	SG	CYS	1	53.070	84.744	4.510 1.00 92.39
	ATOM	5	N	CYS	1	56.229	84.922	2.302 1.00 93.44
	ATOM	6	CA	CYS	1	55.559	84.100	3.317 1.00 98.53
10	ATOM	7	N	SER	2	55.852	82.054	4.483 1.00104.82
	ATOM	8	CA	SER	2	56.093	80.662	4.836 1.00107.58
	ATOM	9	CB	SER	2	57.479	80.140	4.413 1.00109.37
	ATOM	10	OG	SER	2	57.407	78.812	3.938 1.00110.15
	ATOM	11	C	SER	2	55.857	80.530	6.364 1.00108.57
15	ATOM	12	O	SER	2	54.715	80.694	6.799 1.00109.82
	ATOM	13	N	GLN	3	56.875	80.242	7.176 1.00108.52
	ATOM	14	CA	GLN	3	56.662	79.984	8.614 1.00109.56
	ATOM	15	CB	GLN	3	56.614	81.279	9.447 1.00108.23
	ATOM	16	CG	GLN	3	55.323	82.064	9.362 1.00109.80
20	ATOM	17	CD	GLN	3	55.541	83.270	8.494 1.00110.35
	ATOM	18	OE1	GLN	3	56.395	84.110	8.756 1.00110.92
	ATOM	19	NE2	GLN	3	54.863	83.552	7.393 1.00110.59
	ATOM	20	C	GLN	3	55.408	79.096	8.815 1.00110.79
	ATOM	21	O	GLN	3	54.476	79.447	9.544 1.00114.50
25	ATOM	22	N	ASN	4	55.421	77.932	8.143 1.00109.87
	ATOM	23	CA	ASN	4	54.413	76.839	8.139 1.00107.78
	ATOM	24	CB	ASN	4	54.566	75.993	9.396 1.00109.08
	ATOM	25	CG	ASN	4	54.595	76.799	10.690 1.00109.99
	ATOM	26	OD1	ASN	4	55.557	77.515	10.972 1.00109.73
30	ATOM	27	ND2	ASN	4	53.531	76.669	11.485 1.00109.43
	ATOM	28	C	ASN	4	52.969	77.304	7.931 1.00105.15
	ATOM	29	O	ASN	4	52.128	77.145	8.802 1.00103.42
	ATOM	30	N	GLU	5	52.686	77.914	6.753 1.00101.11
	ATOM	31	CA	GLU	5	51.311	78.421	6.424 1.00 95.31
35	ATOM	32	CB	GLU	5	51.080	79.792	7.059 1.00 92.47
	ATOM	33	CG	GLU	5	51.840	80.049	8.350 1.00 86.82
	ATOM	34	CD	GLU	5	52.493	81.416	8.295 1.00 85.55
	ATOM	35	OE1	GLU	5	53.058	81.750	7.240 1.00 85.72
	ATOM	36	OE2	GLU	5	52.425	82.132	9.306 1.00 81.01
40	ATOM	37	C	GLU	5	50.955	78.522	4.919 1.00 92.66
	ATOM	38	O	GLU	5	51.502	77.826	4.081 1.00 94.43
	ATOM	39	N	TYR	6	50.010	79.419	4.638 1.00 85.19
	ATOM	40	CA	TYR	6	49.507	79.660	3.283 1.00 78.14
	ATOM	41	CB	TYR	6	48.348	78.735	2.966 1.00 76.59
45	ATOM	42	CG	TYR	6	47.045	79.118	3.596 1.00 73.94
	ATOM	43	CD1	TYR	6	46.072	79.763	2.818 1.00 73.49
	ATOM	44	CE1	TYR	6	44.851	80.110	3.362 1.00 73.13
	ATOM	45	CD2	TYR	6	46.751	78.849	4.936 1.00 72.45
	ATOM	46	CE2	TYR	6	45.513	79.204	5.493 1.00 72.38
50	ATOM	47	CZ	TYR	6	44.576	79.822	4.701 1.00 74.15
	ATOM	48	OH	TYR	6	43.359	80.179	5.232 1.00 73.32
	ATOM	49	C	TYR	6	49.085	81.109	3.185 1.00 75.43
	ATOM	50	O	TYR	6	48.843	81.763	4.217 1.00 76.93
	ATOM	51	N	PHE	7	48.985	81.630	1.965 1.00 69.11
55	ATOM	52	CA	PHE	7	48.551	83.010	1.788 1.00 62.54
	ATOM	53	CB	PHE	7	49.452	83.703	0.761 1.00 61.14
	ATOM	54	CG	PHE	7	48.993	85.102	0.452 1.00 61.11
	ATOM	55	CD1	PHE	7	49.247	86.116	1.369 1.00 59.90
	ATOM	56	CD2	PHE	7	48.288	85.407	-0.711 1.00 63.23
60	ATOM	57	CE1	PHE	7	48.803	87.418	1.130 1.00 61.45
	ATOM	58	CE2	PHE	7	47.835	86.707	-0.967 1.00 61.15
	ATOM	59	CZ	PHE	7	48.091	87.713	-0.047 1.00 61.89
	ATOM	60	C	PHE	7	47.075	83.081	1.364 1.00 60.79
	ATOM	61	O	PHE	7	46.765	82.958	0.177 1.00 63.10
65	ATOM	62	N	ASP	8	46.171	83.277	2.328 1.00 57.06
	ATOM	63	CA	ASP	8	44.734	83.364	2.039 1.00 52.20

	ATOM	64	CB	ASP	8	43.909	83.364	3.323	1.00	46.52
	ATOM	65	CG	ASP	8	42.412	83.183	3.113	1.00	47.00
	ATOM	66	OD1	ASP	8	41.906	83.676	2.082	1.00	49.00
	ATOM	67	OD2	ASP	8	41.762	82.563	3.980	1.00	48.43
5	ATOM	68	C	ASP	8	44.444	84.638	1.234	1.00	53.02
	ATOM	69	O	ASP	8	44.542	85.753	1.756	1.00	53.25
	ATOM	70	N	SER	9	44.092	84.466	-0.041	1.00	55.34
	ATOM	71	CA	SER	9	43.779	85.600	-0.920	1.00	54.13
	ATOM	72	CB	SER	9	43.600	85.134	-2.368	1.00	54.65
10	ATOM	73	OG	SER	9	44.788	84.534	-2.853	1.00	54.09
	ATOM	74	C	SER	9	42.512	86.338	-0.483	1.00	54.96
	ATOM	75	O	SER	9	42.357	87.541	-0.756	1.00	54.11
	ATOM	76	N	LEU	10	41.604	85.643	0.194	1.00	49.64
	ATOM	77	CA	LEU	10	40.402	86.310	0.650	1.00	46.11
15	ATOM	78	CB	LEU	10	39.422	85.332	1.281	1.00	40.39
	ATOM	79	CG	LEU	10	38.050	85.882	1.606	1.00	34.58
	ATOM	80	CD1	LEU	10	37.296	86.226	0.320	1.00	33.04
	ATOM	81	CD2	LEU	10	37.253	84.896	2.463	1.00	33.82
	ATOM	82	C	LEU	10	40.828	87.397	1.638	1.00	49.82
20	ATOM	83	O	LEU	10	40.364	88.523	1.563	1.00	50.94
	ATOM	84	N	LEU	11	41.714	87.047	2.569	1.00	51.20
	ATOM	85	CA	LEU	11	42.173	88.003	3.575	1.00	51.39
	ATOM	86	CB	LEU	11	42.556	87.272	4.866	1.00	47.82
	ATOM	87	CG	LEU	11	41.502	86.371	5.527	1.00	46.28
25	ATOM	88	CD1	LEU	11	42.050	85.781	6.821	1.00	49.44
	ATOM	89	CD2	LEU	11	40.215	87.136	5.799	1.00	44.53
	ATOM	90	C	LEU	11	43.343	88.890	3.135	1.00	53.87
	ATOM	91	O	LEU	11	43.550	89.965	3.684	1.00	50.27
	ATOM	92	N	HIS	12	44.119	88.430	2.148	1.00	58.57
30	ATOM	93	CA	HIS	12	45.282	89.146	1.666	1.00	64.33
	ATOM	94	CB	HIS	12	44.940	90.578	1.259	1.00	65.68
	ATOM	95	CG	HIS	12	44.008	90.651	0.021	1.00	69.20
	ATOM	96	CD2	HIS	12	43.878	89.825	-1.044	1.00	69.34
	ATOM	97	ND1	HIS	12	43.075	91.651	-0.156	1.00	71.64
35	ATOM	98	CE1	HIS	12	42.403	91.433	-1.277	1.00	72.29
	ATOM	99	NE2	HIS	12	42.872	90.330	-1.835	1.00	70.37
	ATOM	100	C	HIS	12	46.337	89.232	2.761	1.00	66.69
	ATOM	101	O	HIS	12	46.976	90.267	2.979	1.00	67.11
	ATOM	102	N	ALA	13	46.493	88.116	3.467	1.00	69.57
40	ATOM	103	CA	ALA	13	47.428	88.014	4.579	1.00	71.62
	ATOM	104	CB	ALA	13	46.800	88.493	5.866	1.00	69.66
	ATOM	105	C	ALA	13	47.876	86.560	4.706	1.00	73.99
	ATOM	106	O	ALA	13	47.488	85.715	3.891	1.00	75.45
	ATOM	107	N	CYS	14	48.673	86.261	5.730	1.00	75.75
45	ATOM	108	CA	CYS	14	49.168	84.901	5.907	1.00	77.01
	ATOM	109	C	CYS	14	48.598	84.172	7.105	1.00	75.86
	ATOM	110	O	CYS	14	48.755	84.583	8.259	1.00	73.08
	ATOM	111	CB	CYS	14	50.697	84.915	5.940	1.00	80.96
	ATOM	112	SG	CYS	14	51.490	84.507	4.355	1.00	86.58
50	ATOM	113	N	ILE	15	47.924	83.047	6.782	1.00	75.89
	ATOM	114	CA	ILE	15	47.274	82.239	7.800	1.00	77.02
	ATOM	115	CB	ILE	15	45.788	81.988	7.449	1.00	76.68
	ATOM	116	CG2	ILE	15	45.127	81.123	8.518	1.00	76.37
	ATOM	117	CG1	ILE	15	45.022	83.313	7.287	1.00	76.28
55	ATOM	118	CD1	ILE	15	44.901	84.129	8.556	1.00	74.77
	ATOM	119	C	ILE	15	47.977	80.886	8.056	1.00	78.19
	ATOM	120	O	ILE	15	48.359	80.159	7.150	1.00	77.54
	ATOM	121	N	PRO	16	48.092	80.600	9.381	1.00	80.09
	ATOM	122	CD	PRO	16	48.536	81.646	10.332	1.00	79.64
60	ATOM	123	CA	PRO	16	48.668	79.294	9.822	1.00	82.16
	ATOM	124	CB	PRO	16	48.614	79.411	11.335	1.00	80.00
	ATOM	125	CG	PRO	16	48.927	80.852	11.546	1.00	80.17
	ATOM	126	C	PRO	16	48.053	78.098	9.155	1.00	84.28
	ATOM	127	O	PRO	16	46.835	77.942	9.214	1.00	84.12
65	ATOM	128	N	CYS	17	48.845	77.238	8.507	1.00	87.37
	ATOM	129	CA	CYS	17	48.281	76.069	7.849	1.00	88.33

	ATOM	130	CB	CYS	17	49.392	75.138	7.373	1.00	86.90
	ATOM	131	SG	CYS	17	50.200	75.655	5.829	1.00	84.56
	ATOM	132	C	CYS	17	47.322	75.310	8.773	1.00	89.64
	ATOM	133	O	CYS	17	46.266	74.824	8.335	1.00	89.77
5	ATOM	134	N	GLN	18	47.705	75.210	10.047	1.00	91.08
	ATOM	135	CA	GLN	18	46.895	74.560	11.035	1.00	91.16
	ATOM	136	CB	GLN	18	47.256	74.999	12.455	1.00	92.68
	ATOM	137	CG	GLN	18	48.759	75.135	12.671	1.00	98.31
	ATOM	138	CD	GLN	18	49.155	76.053	13.833	1.00	101.37
10	ATOM	139	OE1	GLN	18	48.983	77.264	13.765	1.00	102.74
	ATOM	140	NE2	GLN	18	49.690	75.669	14.977	1.00	103.76
	ATOM	141	C	GLN	18	45.426	74.888	10.757	1.00	89.18
	ATOM	142	O	GLN	18	44.587	74.000	10.590	1.00	89.64
	ATOM	143	N	LEU	19	45.156	76.194	10.725	1.00	87.56
15	ATOM	144	CA	LEU	19	43.826	76.741	10.525	1.00	85.33
	ATOM	145	CB	LEU	19	43.909	78.227	10.325	1.00	79.98
	ATOM	146	CG	LEU	19	43.515	79.024	11.551	1.00	77.91
	ATOM	147	CD1	LEU	19	43.974	80.468	11.423	1.00	78.13
	ATOM	148	CD2	LEU	19	42.016	78.957	11.782	1.00	72.50
20	ATOM	149	C	LEU	19	43.039	76.120	9.383	1.00	86.05
	ATOM	150	O	LEU	19	41.849	75.808	9.548	1.00	88.06
	ATOM	151	N	ARG	20	43.664	75.926	8.247	1.00	85.11
	ATOM	152	CA	ARG	20	42.932	75.375	7.101	1.00	83.33
	ATOM	153	CB	ARG	20	43.613	75.781	5.797	1.00	78.93
25	ATOM	154	CG	ARG	20	42.913	76.903	5.036	1.00	73.68
	ATOM	155	CD	ARG	20	41.561	76.467	4.501	1.00	67.23
	ATOM	156	NE	ARG	20	41.531	76.454	3.038	1.00	63.84
	ATOM	157	CZ	ARG	20	41.939	77.468	2.271	1.00	64.39
	ATOM	158	NH1	ARG	20	42.407	78.573	2.843	1.00	66.99
30	ATOM	159	NH2	ARG	20	41.870	77.370	0.953	1.00	60.63
	ATOM	160	C	ARG	20	42.753	73.857	7.163	1.00	84.21
	ATOM	161	O	ARG	20	42.474	73.239	6.136	1.00	86.28
	ATOM	162	N	CYS	21	42.906	73.259	8.336	1.00	85.02
	ATOM	163	CA	CYS	21	42.705	71.817	8.422	1.00	83.51
35	ATOM	164	CB	CYS	21	43.938	71.165	9.065	1.00	81.05
	ATOM	165	SG	CYS	21	45.500	71.475	8.182	1.00	73.08
	ATOM	166	C	CYS	21	41.396	71.433	9.158	1.00	84.26
	ATOM	167	O	CYS	21	41.465	70.896	10.254	1.00	84.48
	ATOM	168	N	SER	22	40.182	71.725	8.569	1.00	84.61
40	ATOM	169	CA	SER	22	38.832	71.414	9.133	1.00	85.07
	ATOM	170	C	SER	22	38.140	70.350	8.292	1.00	85.65
	ATOM	171	O	SER	22	36.918	70.229	8.263	1.00	86.77
	ATOM	172	CB	SER	22	37.930	72.655	9.167	1.00	20.00
	ATOM	173	OG	SER	22	37.249	72.817	7.938	1.00	20.00
45	ATOM	174	N	SER	23	38.988	69.604	7.637	1.00	84.19
	ATOM	175	CA	SER	23	38.550	68.492	6.825	1.00	82.44
	ATOM	176	C	SER	23	38.520	68.797	5.324	1.00	84.32
	ATOM	177	O	SER	23	39.572	68.732	4.686	1.00	81.24
	ATOM	178	CB	SER	23	37.198	68.014	7.355	1.00	78.00
50	ATOM	179	OG	SER	23	36.352	67.586	6.306	1.00	20.00
	ATOM	180	N	ASN	24	37.396	69.118	4.748	1.00	87.06
	ATOM	181	CA	ASN	24	37.493	69.392	3.329	1.00	88.17
	ATOM	182	C	ASN	24	38.419	70.599	3.117	1.00	88.45
	ATOM	183	O	ASN	24	38.564	71.058	1.997	1.00	84.89
55	ATOM	184	CB	ASN	24	36.155	69.762	2.728	0.00	86.84
	ATOM	185	CG	ASN	24	35.102	68.752	3.008	0.00	85.55
	ATOM	186	OD1	ASN	24	35.407	67.577	3.196	1.00	20.00
	ATOM	187	ND2	ASN	24	33.847	69.179	3.037	1.00	20.00
	ATOM	188	N	THR	25	39.039	71.113	4.212	1.00	89.49
60	ATOM	189	CA	THR	25	39.798	72.372	4.066	1.00	91.86
	ATOM	190	C	THR	25	41.364	72.367	4.013	1.00	95.75
	ATOM	191	O	THR	25	42.023	73.358	4.372	1.00	96.59
	ATOM	192	CB	THR	25	39.278	73.337	5.109	0.00	91.23
	ATOM	193	OG1	THR	25	37.864	73.473	4.957	1.00	20.00
65	ATOM	194	CG2	THR	25	39.955	74.682	4.956	1.00	20.00
	ATOM	195	N	PRO	26	41.948	71.231	3.567	1.00	98.41

	ATOM	196	CA	PRO	26	43.472	71.097	3.243	1.00	96.39
	ATOM	197	C	PRO	26	44.180	71.792	1.908	1.00	96.88
	ATOM	198	O	PRO	26	44.156	71.126	0.869	1.00	94.52
	ATOM	199	CB	PRO	26	43.614	69.587	3.240	0.00	95.29
5	ATOM	200	CG	PRO	26	42.701	69.125	4.336	0.00	92.34
	ATOM	201	CD	PRO	26	41.651	70.172	4.530	0.00	89.50
	ATOM	202	N	PRO	27	44.771	73.090	1.891	1.00	97.72
	ATOM	203	CA	PRO	27	45.542	73.739	0.683	1.00	98.94
	ATOM	204	C	PRO	27	46.867	73.120	0.061	1.00	100.71
10	ATOM	205	O	PRO	27	47.461	72.259	0.705	1.00	100.56
	ATOM	206	CB	PRO	27	45.682	75.172	1.144	0.00	97.33
	ATOM	207	CG	PRO	27	44.384	75.413	1.843	0.00	94.51
	ATOM	208	CD	PRO	27	43.849	74.118	2.337	0.00	92.58
	ATOM	209	N	LEU	28	47.373	73.510	-1.176	1.00	15.00
15	ATOM	210	CA	LEU	28	48.722	72.961	-1.538	1.00	15.00
	ATOM	211	C	LEU	28	49.495	73.347	-0.280	1.00	15.00
	ATOM	212	O	LEU	28	48.886	73.594	0.762	1.00	15.00
	ATOM	213	CB	LEU	28	49.351	73.594	-2.827	1.00	15.00
	ATOM	214	CG	LEU	28	49.667	72.656	-4.035	1.00	15.00
20	ATOM	215	CD1	LEU	28	51.160	72.561	-4.245	1.00	15.00
	ATOM	216	CD2	LEU	28	49.055	71.273	-3.819	1.00	15.00
	ATOM	217	N	THR	29	50.810	73.427	-0.394	1.00	15.00
	ATOM	218	CA	THR	29	51.603	73.881	0.742	1.00	15.00
	ATOM	219	C	THR	29	50.837	73.676	2.045	1.00	15.00
25	ATOM	220	O	THR	29	51.100	74.340	3.035	1.00	15.00
	ATOM	221	CB	THR	29	51.994	75.351	0.552	1.00	15.00
	ATOM	222	OG1	THR	29	53.334	75.417	0.075	1.00	20.00
	ATOM	223	CG2	THR	29	51.868	76.095	1.875	1.00	20.00
	ATOM	224	N	CYS	30	49.893	72.721	1.999	1.00	15.00
30	ATOM	225	CA	CYS	30	49.076	72.477	3.171	1.00	15.00
	ATOM	226	C	CYS	30	48.847	70.983	3.363	1.00	15.00
	ATOM	227	O	CYS	30	48.691	70.530	4.486	1.00	15.00
	ATOM	228	CB	CYS	30	47.784	73.278	3.096	1.00	15.00
	ATOM	229	SG	CYS	30	47.974	75.066	3.376	1.00	20.00
35	ATOM	230	N	GLN	31	48.840	70.234	2.260	1.00	15.00
	ATOM	231	CA	GLN	31	48.674	68.804	2.368	1.00	15.00
	ATOM	232	C	GLN	31	49.772	68.179	3.232	1.00	15.00
	ATOM	233	O	GLN	31	49.507	67.404	4.157	1.00	15.00
	ATOM	234	CB	GLN	31	48.647	68.162	1.004	1.00	15.00
40	ATOM	235	CG	GLN	31	47.241	68.177	0.418	1.00	20.00
	ATOM	236	CD	GLN	31	47.150	67.446	-0.907	1.00	20.00
	ATOM	237	OE1	GLN	31	48.016	66.639	-1.243	1.00	20.00
	ATOM	238	NE2	GLN	31	46.188	67.576	-1.814	1.00	20.00
	ATOM	239	N	ARG	32	50.994	68.524	2.934	1.00	15.00
45	ATOM	240	CA	ARG	32	52.145	68.063	3.682	1.00	15.00
	ATOM	241	C	ARG	32	52.047	68.431	5.159	1.00	15.00
	ATOM	242	O	ARG	32	52.393	67.632	6.046	1.00	15.00
	ATOM	243	CB	ARG	32	53.438	68.594	3.028	1.00	15.00
	ATOM	244	CG	ARG	32	53.669	68.122	1.589	1.00	15.00
50	ATOM	245	CD	ARG	32	55.008	68.637	1.068	1.00	15.00
	ATOM	246	NE	ARG	32	55.247	68.279	-0.325	1.00	20.00
	ATOM	247	CZ	ARG	32	56.365	68.561	-0.983	1.00	20.00
	ATOM	248	NH1	ARG	32	57.349	69.206	-0.370	1.00	20.00
	ATOM	249	NH2	ARG	32	56.502	68.197	-2.251	1.00	20.00
55	ATOM	250	N	TYR	33	51.552	69.644	5.393	1.00	15.00
	ATOM	251	CA	TYR	33	51.375	70.142	6.746	1.00	15.00
	ATOM	252	C	TYR	33	50.297	69.346	7.534	1.00	15.00
	ATOM	253	O	TYR	33	50.556	68.917	8.651	1.00	15.00
	ATOM	254	CB	TYR	33	51.039	71.597	6.668	1.00	15.00
60	ATOM	255	CG	TYR	33	52.090	72.400	5.924	1.00	20.00
	ATOM	256	CD1	TYR	33	52.012	72.626	4.557	1.00	20.00
	ATOM	257	CD2	TYR	33	53.156	72.956	6.619	1.00	20.00
	ATOM	258	CE1	TYR	33	52.963	73.383	3.903	1.00	20.00
	ATOM	259	CE2	TYR	33	54.111	73.717	5.974	1.00	20.00
65	ATOM	260	CZ	TYR	33	54.011	73.928	4.616	1.00	20.00
	ATOM	261	OH	TYR	33	54.960	74.685	3.969	1.00	20.00

	ATOM	262	N	CYS	34	49.103	69.150	6.949	1.00	15.00
	ATOM	263	CA	CYS	34	48.041	68.392	7.593	1.00	15.00
	ATOM	264	C	CYS	34	48.582	67.001	8.007	1.00	15.00
	ATOM	265	O	CYS	34	48.214	66.476	9.059	1.00	15.00
5	ATOM	266	CB	CYS	34	46.819	68.294	6.687	1.00	15.00
	ATOM	267	SG	CYS	34	45.969	69.873	6.376	1.00	20.00
	ATOM	268	N	GLN	35	49.451	66.440	7.171	1.00	15.00
	ATOM	269	CA	GLN	35	50.129	65.140	7.349	1.00	15.00
	ATOM	270	CB	GLN	35	50.831	64.798	6.033	1.00	15.00
10	ATOM	271	CG	GLN	35	51.219	63.354	5.845	1.00	15.00
	ATOM	272	CD	GLN	35	50.065	62.446	6.154	1.00	15.00
	ATOM	273	OE1	GLN	35	48.980	62.608	5.599	1.00	15.00
	ATOM	274	NE2	GLN	35	50.068	61.423	6.984	1.00	15.00
	ATOM	275	C	GLN	35	51.177	65.167	8.458	1.00	15.00
15	ATOM	276	O	GLN	35	51.223	64.269	9.304	1.00	15.00
	ATOM	277	N	ALA	36	51.986	66.220	8.430	1.00	15.00
	ATOM	278	CA	ALA	36	53.023	66.479	9.431	1.00	15.00
	ATOM	279	C	ALA	36	52.377	66.460	10.829	1.00	15.00
	ATOM	280	O	ALA	36	52.873	65.784	11.733	1.00	15.00
20	ATOM	281	CB	ALA	36	53.705	67.801	9.181	1.00	15.00
	ATOM	282	N	SER	37	51.278	67.240	10.974	1.00	15.00
	ATOM	283	CA	SER	37	50.625	67.313	12.302	1.00	15.00
	ATOM	284	C	SER	37	49.966	65.989	12.610	1.00	15.00
	ATOM	285	O	SER	37	49.999	65.515	13.759	1.00	15.00
25	ATOM	286	CB	SER	37	49.658	68.505	12.387	1.00	15.00
	ATOM	287	OG	SER	37	48.778	68.549	11.278	1.00	20.00
	ATOM	288	N	VAL	38	49.360	65.383	11.585	1.00	15.00
	ATOM	289	CA	VAL	38	48.739	64.098	11.738	1.00	15.00
	ATOM	290	C	VAL	38	49.709	63.065	12.324	1.00	15.00
30	ATOM	291	O	VAL	38	49.414	62.378	13.320	1.00	15.00
	ATOM	292	CB	VAL	38	48.222	63.577	10.364	1.00	15.00
	ATOM	293	CG1	VAL	38	47.829	62.102	10.442	1.00	20.00
	ATOM	294	CG2	VAL	38	47.027	64.406	9.900	1.00	20.00
	ATOM	295	N	THR	39	50.871	62.946	11.659	1.00	15.00
35	ATOM	296	CA	THR	39	51.890	61.987	12.079	1.00	15.00
	ATOM	297	C	THR	39	52.061	62.026	13.590	1.00	15.00
	ATOM	298	O	THR	39	52.805	61.220	14.158	1.00	15.00
	ATOM	299	CB	THR	39	53.227	62.320	11.397	1.00	15.00
	ATOM	300	OG1	THR	39	53.089	62.173	9.973	1.00	20.00
40	ATOM	301	CG2	THR	39	54.309	61.403	11.917	1.00	20.00
	END					134.002	98.540	-12.573	0.00	0.00

TABLE 17

	111								
	ATOM	1	C	CYS	1	22.235	102.632	33.513	1.00101.54
5	ATOM	2	O	CYS	1	22.654	103.749	33.806	1.00102.94
	ATOM	3	CB	CYS	1	21.597	102.068	31.160	1.00 96.91
	ATOM	4	SG	CYS	1	21.152	100.396	30.593	1.00 92.39
	ATOM	5	N	CYS	1	20.205	103.617	32.495	1.00 93.44
	ATOM	6	CA	CYS	1	21.069	102.432	32.563	1.00 98.53
10	ATOM	7	N	SER	2	22.733	101.473	33.957	1.00104.82
	ATOM	8	CA	SER	2	23.907	101.244	34.788	1.00107.58
	ATOM	9	CB	SER	2	23.983	102.155	36.027	1.00109.37
	ATOM	10	OG	SER	2	25.307	102.579	36.276	1.00110.15
	ATOM	11	C	SER	2	23.916	99.739	35.163	1.00108.57
15	ATOM	12	O	SER	2	24.091	98.910	34.267	1.00109.82
	ATOM	13	N	GLN	3	23.740	99.364	36.431	1.00108.52
	ATOM	14	CA	GLN	3	23.871	97.952	36.840	1.00109.56
	ATOM	15	CB	GLN	3	22.573	97.151	36.625	1.00108.23
	ATOM	16	CG	GLN	3	22.284	96.745	35.196	1.00109.80
20	ATOM	17	CD	GLN	3	21.195	97.623	34.650	1.00110.35
	ATOM	18	OE1	GLN	3	20.091	97.690	35.179	1.00110.92
	ATOM	19	NE2	GLN	3	21.298	98.392	33.578	1.00110.59
	ATOM	20	C	GLN	3	25.095	97.307	36.143	1.00110.79
	ATOM	21	O	GLN	3	24.988	96.281	35.464	1.00114.50
25	ATOM	22	N	ASN	4	26.262	97.944	36.337	1.00109.87
	ATOM	23	CA	ASN	4	27.621	97.582	35.855	1.00107.78
	ATOM	24	CB	ASN	4	28.201	96.478	36.728	1.00109.08
	ATOM	25	CG	ASN	4	27.275	95.281	36.916	1.00109.99
	ATOM	26	OD1	ASN	4	26.248	95.370	37.591	1.00109.73
30	ATOM	27	ND2	ASN	4	27.653	94.149	36.318	1.00109.43
	ATOM	28	C	ASN	4	27.698	97.236	34.365	1.00105.15
	ATOM	29	O	ASN	4	28.019	96.117	33.997	1.00103.42
	ATOM	30	N	GLU	5	27.373	98.221	33.490	1.00101.11
	ATOM	31	CA	GLU	5	27.403	98.013	32.003	1.00 95.31
35	ATOM	32	CB	GLU	5	26.120	97.327	31.533	1.00 92.47
	ATOM	33	CG	GLU	5	25.461	96.408	32.549	1.00 86.82
	ATOM	34	CD	GLU	5	23.973	96.691	32.605	1.00 85.55
	ATOM	35	OE1	GLU	5	23.605	97.878	32.607	1.00 85.72
	ATOM	36	OE2	GLU	5	23.200	95.722	32.642	1.00 81.01
40	ATOM	37	C	GLU	5	27.618	99.278	31.134	1.00 92.66
	ATOM	38	O	GLU	5	28.190	100.264	31.565	1.00 94.43
	ATOM	39	N	TYR	6	27.133	99.182	29.896	1.00 85.19
	ATOM	40	CA	TYR	6	27.248	100.252	28.902	1.00 78.14
	ATOM	41	CB	TYR	6	28.541	100.123	28.119	1.00 76.59
45	ATOM	42	CG	TYR	6	28.541	99.052	27.073	1.00 73.94
	ATOM	43	CD1	TYR	6	28.364	99.409	25.728	1.00 73.49
	ATOM	44	CE1	TYR	6	28.381	98.449	24.736	1.00 73.13
	ATOM	45	CD2	TYR	6	28.721	97.701	27.382	1.00 72.45
	ATOM	46	CE2	TYR	6	28.735	96.722	26.377	1.00 72.38
50	ATOM	47	CZ	TYR	6	28.572	97.106	25.068	1.00 74.15
	ATOM	48	OH	TYR	6	28.580	96.159	24.071	1.00 73.32
	ATOM	49	C	TYR	6	26.050	100.176	27.983	1.00 75.43
	ATOM	50	O	TYR	6	25.390	99.123	27.899	1.00 76.93
	ATOM	51	N	PHE	7	25.752	101.268	27.284	1.00 69.11
55	ATOM	52	CA	PHE	7	24.631	101.261	26.352	1.00 62.54
	ATOM	53	CB	PHE	7	23.812	102.544	26.526	1.00 61.14
	ATOM	54	CG	PHE	7	22.699	102.650	25.519	1.00 61.11
	ATOM	55	CD1	PHE	7	21.551	101.886	25.695	1.00 59.90
	ATOM	56	CD2	PHE	7	22.798	103.467	24.394	1.00 63.23
60	ATOM	57	CE1	PHE	7	20.515	101.933	24.761	1.00 61.45
	ATOM	58	CE2	PHE	7	21.769	103.527	23.446	1.00 61.15
	ATOM	59	CZ	PHE	7	20.627	102.760	23.628	1.00 61.89
	ATOM	60	C	PHE	7	25.114	101.107	24.901	1.00 60.79
	ATOM	61	O	PHE	7	25.483	102.096	24.262	1.00 63.10
65	ATOM	62	N	ASP	8	25.112	99.875	24.383	1.00 57.06
	ATOM	63	CA	ASP	8	25.550	99.610	23.007	1.00 52.20

	ATOM	64	CB	ASP	8	25.664	98.112	22.740	1.00	46.52
	ATOM	65	CG	ASP	8	26.362	97.753	21.436	1.00	47.00
	ATOM	66	OD1	ASP	8	26.202	98.520	20.461	1.00	49.00
	ATOM	67	OD2	ASP	8	27.049	96.712	21.397	1.00	48.43
5	ATOM	68	C	ASP	8	24.560	100.241	22.018	1.00	53.02
	ATOM	69	O	ASP	8	23.420	99.784	21.886	1.00	53.25
	ATOM	70	N	SER	9	24.999	101.296	21.331	1.00	55.34
	ATOM	71	CA	SER	9	24.157	101.988	20.347	1.00	54.13
	ATOM	72	CB	SER	9	24.836	103.270	19.854	1.00	54.65
10	ATOM	73	OG	SER	9	25.058	104.165	20.929	1.00	54.09
	ATOM	74	C	SER	9	23.838	101.107	19.137	1.00	54.96
	ATOM	75	O	SER	9	22.802	101.294	18.477	1.00	54.11
	ATOM	76	N	LEU	10	24.706	100.146	18.836	1.00	49.64
	ATOM	77	CA	LEU	10	24.429	99.272	17.715	1.00	46.11
15	ATOM	78	CB	LEU	10	25.591	98.330	17.437	1.00	40.39
	ATOM	79	CG	LEU	10	25.496	97.516	16.165	1.00	34.58
	ATOM	80	CD1	LEU	10	25.591	98.428	14.940	1.00	33.04
	ATOM	81	CD2	LEU	10	26.576	96.432	16.127	1.00	33.82
	ATOM	82	C	LEU	10	23.146	98.505	18.038	1.00	49.82
20	ATOM	83	O	LEU	10	22.260	98.394	17.207	1.00	50.94
	ATOM	84	N	LEU	11	23.058	97.971	19.256	1.00	51.20
	ATOM	85	CA	LEU	11	21.884	97.200	19.661	1.00	51.39
	ATOM	86	CB	LEU	11	22.275	96.148	20.705	1.00	47.82
	ATOM	87	CG	LEU	11	23.386	95.151	20.346	1.00	46.28
25	ATOM	88	CD1	LEU	11	23.589	94.156	21.483	1.00	49.44
	ATOM	89	CD2	LEU	11	23.069	94.415	19.052	1.00	44.53
	ATOM	90	C	LEU	11	20.719	98.036	20.203	1.00	53.87
	ATOM	91	O	LEU	11	19.577	97.595	20.188	1.00	50.27
	ATOM	92	N	HIS	12	21.013	99.244	20.696	1.00	58.57
30	ATOM	93	CA	HIS	12	20.015	100.117	21.279	1.00	64.33
	ATOM	94	CB	HIS	12	18.845	100.358	20.327	1.00	65.68
	ATOM	95	CG	HIS	12	19.246	101.161	19.062	1.00	69.20
	ATOM	96	CD2	HIS	12	20.196	102.109	18.875	1.00	69.34
	ATOM	97	ND1	HIS	12	18.648	100.973	17.834	1.00	71.64
35	ATOM	98	CE1	HIS	12	19.219	101.766	16.940	1.00	72.29
	ATOM	99	NE2	HIS	12	20.162	102.466	17.547	1.00	70.37
	ATOM	100	C	HIS	12	19.443	99.491	22.543	1.00	66.69
	ATOM	101	O	HIS	12	18.235	99.518	22.801	1.00	67.11
	ATOM	102	N	ALA	13	20.343	98.901	23.324	1.00	69.57
40	ATOM	103	CA	ALA	13	19.984	98.216	24.558	1.00	71.62
	ATOM	104	CB	ALA	13	19.584	96.784	24.291	1.00	69.66
	ATOM	105	C	ALA	13	21.174	98.275	25.512	1.00	73.99
	ATOM	106	O	ALA	13	22.197	98.894	25.195	1.00	75.45
	ATOM	107	N	CYS	14	21.057	97.622	26.666	1.00	75.75
45	ATOM	108	CA	CYS	14	22.138	97.651	27.644	1.00	77.01
	ATOM	109	C	CYS	14	22.858	96.332	27.829	1.00	75.86
	ATOM	110	O	CYS	14	22.276	95.316	28.220	1.00	73.08
	ATOM	111	CB	CYS	14	21.607	98.187	28.975	1.00	80.96
	ATOM	112	SG	CYS	14	21.923	99.956	29.255	1.00	86.58
50	ATOM	113	N	ILE	15	24.175	96.391	27.536	1.00	75.89
	ATOM	114	CA	ILE	15	25.019	95.210	27.618	1.00	77.02
	ATOM	115	CB	ILE	15	25.796	94.987	26.299	1.00	76.68
	ATOM	116	CG2	ILE	15	26.690	93.756	26.409	1.00	76.37
	ATOM	117	CG1	ILE	15	24.837	94.844	25.104	1.00	76.28
55	ATOM	118	CD1	ILE	15	23.955	93.614	25.149	1.00	74.77
	ATOM	119	C	ILE	15	26.014	95.243	28.801	1.00	78.19
	ATOM	120	O	ILE	15	26.679	96.231	29.077	1.00	77.54
	ATOM	121	N	PRO	16	26.074	94.057	29.465	1.00	80.09
	ATOM	122	CD	PRO	16	24.829	93.331	29.806	1.00	79.64
60	ATOM	123	CA	PRO	16	27.044	93.871	30.586	1.00	82.16
	ATOM	124	CB	PRO	16	26.762	92.445	31.025	1.00	80.00
	ATOM	125	CG	PRO	16	25.285	92.355	30.853	1.00	80.17
	ATOM	126	C	PRO	16	28.451	94.271	30.250	1.00	84.28
	ATOM	127	O	PRO	16	28.998	93.766	29.271	1.00	84.12
65	ATOM	128	N	CYS	17	29.071	95.173	31.018	1.00	87.37
	ATOM	129	CA	CYS	17	30.434	95.583	30.719	1.00	88.33

	ATOM	130	CB	CYS	17	30.991	96.444	31.849	1.00	86.90
	ATOM	131	SG	CYS	17	30.433	98.174	31.825	1.00	84.56
	ATOM	132	C	CYS	17	31.348	94.375	30.483	1.00	89.64
	ATOM	133	O	CYS	17	32.211	94.394	29.590	1.00	89.77
5	ATOM	134	N	GLN	18	31.151	93.335	31.294	1.00	91.08
	ATOM	135	CA	GLN	18	31.905	92.122	31.171	1.00	91.16
	ATOM	136	CB	GLN	18	31.194	90.934	31.821	1.00	92.68
	ATOM	137	CG	GLN	18	30.536	91.290	33.149	1.00	98.31
	ATOM	138	CD	GLN	18	29.399	90.350	33.568	1.00	101.37
10	ATOM	139	OE1	GLN	18	28.335	90.341	32.961	1.00	102.74
	ATOM	140	NE2	GLN	18	29.433	89.489	34.568	1.00	103.76
	ATOM	141	C	GLN	18	32.127	91.833	29.684	1.00	89.18
	ATOM	142	O	GLN	18	33.258	91.684	29.217	1.00	89.64
	ATOM	143	N	LEU	19	31.002	91.753	28.971	1.00	87.56
15	ATOM	144	CA	LEU	19	30.963	91.442	27.553	1.00	85.33
	ATOM	145	CB	LEU	19	29.573	91.647	27.023	1.00	79.98
	ATOM	146	CG	LEU	19	28.806	90.357	26.823	1.00	77.91
	ATOM	147	CD1	LEU	19	27.321	90.635	26.658	1.00	78.13
	ATOM	148	CD2	LEU	19	29.343	89.587	25.629	1.00	72.50
20	ATOM	149	C	LEU	19	31.951	92.215	26.696	1.00	86.05
	ATOM	150	O	LEU	19	32.621	91.623	25.835	1.00	88.06
	ATOM	151	N	ARG	20	32.066	93.503	26.911	1.00	85.11
	ATOM	152	CA	ARG	20	32.972	94.300	26.075	1.00	83.33
	ATOM	153	CB	ARG	20	32.529	95.761	26.065	1.00	78.93
25	ATOM	154	CG	ARG	20	31.813	96.199	24.791	1.00	73.68
	ATOM	155	CD	ARG	20	32.741	96.198	23.590	1.00	67.23
	ATOM	156	NE	ARG	20	32.948	97.546	23.059	1.00	63.84
	ATOM	157	CZ	ARG	20	31.962	98.402	22.781	1.00	64.39
	ATOM	158	NH1	ARG	20	30.701	98.036	22.990	1.00	66.99
30	ATOM	159	NH2	ARG	20	32.243	99.601	22.297	1.00	60.63
	ATOM	160	C	ARG	20	34.441	94.187	26.486	1.00	84.21
	ATOM	161	O	ARG	20	35.241	95.042	26.108	1.00	86.28
	ATOM	162	N	CYS	21	34.799	93.159	27.242	1.00	85.02
	ATOM	163	CA	CYS	21	36.202	93.015	27.615	1.00	83.51
35	ATOM	164	CB	CYS	21	36.315	92.880	29.140	1.00	81.05
	ATOM	165	SG	CYS	21	35.614	94.277	30.074	1.00	73.08
	ATOM	166	C	CYS	21	36.906	91.849	26.875	1.00	84.26
	ATOM	167	O	CYS	21	37.246	90.861	27.510	1.00	84.48
	ATOM	168	N	SER	22	37.116	91.944	25.514	1.00	84.61
40	ATOM	169	CA	SER	22	37.788	90.922	24.653	1.00	85.07
	ATOM	170	C	SER	22	39.119	91.454	24.142	1.00	85.65
	ATOM	171	O	SER	22	39.646	91.029	23.117	1.00	86.77
	ATOM	172	CB	SER	22	36.927	90.547	23.439	1.00	20.00
	ATOM	173	OG	SER	22	37.160	91.434	22.362	1.00	20.00
45	ATOM	174	N	SER	23	39.614	92.382	24.915	1.00	84.19
	ATOM	175	CA	SER	23	40.902	92.982	24.652	1.00	82.44
	ATOM	176	C	SER	23	40.817	94.363	23.993	1.00	84.32
	ATOM	177	O	SER	23	40.606	95.346	24.705	1.00	81.24
	ATOM	178	CB	SER	23	41.735	91.992	23.837	1.00	78.00
50	ATOM	179	OG	SER	23	42.551	92.656	22.892	1.00	20.00
	ATOM	180	N	ASN	24	40.967	94.479	22.704	1.00	87.06
	ATOM	181	CA	ASN	24	40.859	95.831	22.195	1.00	88.17
	ATOM	182	C	ASN	24	39.448	96.362	22.490	1.00	88.45
	ATOM	183	O	ASN	24	39.113	97.452	22.061	1.00	84.89
55	ATOM	184	CB	ASN	24	41.038	95.890	20.694	0.00	86.84
	ATOM	185	CG	ASN	24	42.299	95.247	20.242	0.00	85.55
	ATOM	186	OD1	ASN	24	43.270	95.194	20.993	1.00	20.00
	ATOM	187	ND2	ASN	24	42.318	94.751	19.012	1.00	20.00
	ATOM	188	N	THR	25	38.622	95.571	23.225	1.00	89.49
60	ATOM	189	CA	THR	25	37.210	95.979	23.380	1.00	91.86
	ATOM	190	C	THR	25	36.696	96.609	24.719	1.00	95.75
	ATOM	191	O	THR	25	35.505	96.512	25.060	1.00	96.59
	ATOM	192	CB	THR	25	36.352	94.810	22.947	0.00	91.23
	ATOM	193	OG1	THR	25	36.719	94.426	21.621	1.00	20.00
65	ATOM	194	CG2	THR	25	34.889	95.194	22.998	1.00	20.00
	ATOM	195	N	PRO	26	37.617	97.248	25.477	1.00	98.41

	ATOM	196	CA	PRO	26	37.272	98.115	26.732	1.00	96.39
	ATOM	197	C	PRO	26	36.555	99.612	26.630	1.00	96.88
	ATOM	198	O	PRO	26	37.316	100.573	26.486	1.00	94.52
	ATOM	199	CB	PRO	26	38.634	98.181	27.395	0.00	95.29
5	ATOM	200	CG	PRO	26	39.233	96.828	27.152	0.00	92.34
	ATOM	201	CD	PRO	26	38.583	96.251	25.935	0.00	89.50
	ATOM	202	N	PRO	27	35.147	99.837	26.671	1.00	97.72
	ATOM	203	CA	PRO	27	34.435	101.240	26.685	1.00	98.94
	ATOM	204	C	PRO	27	34.647	102.314	27.837	1.00	100.71
10	ATOM	205	O	PRO	27	35.170	101.942	28.885	1.00	100.56
	ATOM	206	CB	PRO	27	32.992	100.853	26.453	0.00	97.33
	ATOM	207	CG	PRO	27	33.114	99.721	25.486	0.00	94.51
	ATOM	208	CD	PRO	27	34.440	99.074	25.659	0.00	92.58
	ATOM	209	N	LEU	28	34.270	103.647	27.705	1.00	15.00
15	ATOM	210	CA	LEU	28	34.376	104.488	28.943	1.00	15.00
	ATOM	211	C	LEU	28	33.597	103.603	29.913	1.00	15.00
	ATOM	212	O	LEU	28	33.439	102.408	29.660	1.00	15.00
	ATOM	213	CB	LEU	28	33.737	105.913	28.812	1.00	15.00
	ATOM	214	CG	LEU	28	34.659	107.159	29.000	1.00	15.00
20	ATOM	215	CD1	LEU	28	34.273	107.908	30.254	1.00	15.00
	ATOM	216	CD2	LEU	28	36.128	106.742	29.042	1.00	15.00
	ATOM	217	N	THR	29	33.095	104.196	30.983	1.00	15.00
	ATOM	218	CA	THR	29	32.262	103.431	31.903	1.00	15.00
	ATOM	219	C	THR	29	32.545	101.939	31.768	1.00	15.00
25	ATOM	220	O	THR	29	31.712	101.112	32.102	1.00	15.00
	ATOM	221	CB	THR	29	30.782	103.742	31.648	1.00	15.00
	ATOM	222	OG1	THR	29	30.331	104.682	32.619	1.00	20.00
	ATOM	223	CG2	THR	29	29.962	102.461	31.734	1.00	20.00
	ATOM	224	N	CYS	30	33.759	101.639	31.276	1.00	15.00
30	ATOM	225	CA	CYS	30	34.113	100.249	31.065	1.00	15.00
	ATOM	226	C	CYS	30	35.560	99.997	31.469	1.00	15.00
	ATOM	227	O	CYS	30	35.893	98.900	31.888	1.00	15.00
	ATOM	228	CB	CYS	30	33.808	99.834	29.633	1.00	15.00
	ATOM	229	SG	CYS	30	32.039	99.631	29.254	1.00	20.00
35	ATOM	230	N	GLN	31	36.401	101.024	31.347	1.00	15.00
	ATOM	231	CA	GLN	31	37.778	100.873	31.754	1.00	15.00
	ATOM	232	C	GLN	31	37.883	100.482	33.230	1.00	15.00
	ATOM	233	O	GLN	31	38.579	99.531	33.600	1.00	15.00
	ATOM	234	CB	GLN	31	38.559	102.134	31.485	1.00	15.00
40	ATOM	235	CG	GLN	31	39.091	102.157	30.058	1.00	20.00
	ATOM	236	CD	GLN	31	39.971	103.359	29.779	1.00	20.00
	ATOM	237	OE1	GLN	31	40.477	103.998	30.701	1.00	20.00
	ATOM	238	NE2	GLN	31	40.287	103.844	28.583	1.00	20.00
	ATOM	239	N	ARG	32	37.189	101.209	34.060	1.00	15.00
45	ATOM	240	CA	ARG	32	37.139	100.945	35.483	1.00	15.00
	ATOM	241	C	ARG	32	36.641	99.534	35.781	1.00	15.00
	ATOM	242	O	ARG	32	37.159	98.845	36.677	1.00	15.00
	ATOM	243	CB	ARG	32	36.292	102.028	36.184	1.00	15.00
	ATOM	244	CG	ARG	32	36.837	103.454	36.052	1.00	15.00
50	ATOM	245	CD	ARG	32	35.973	104.430	36.845	1.00	15.00
	ATOM	246	NE	ARG	32	36.403	105.815	36.695	1.00	20.00
	ATOM	247	CZ	ARG	32	35.848	106.839	37.333	1.00	20.00
	ATOM	248	NH1	ARG	32	34.838	106.629	38.168	1.00	20.00
	ATOM	249	NH2	ARG	32	36.303	108.069	37.140	1.00	20.00
55	ATOM	250	N	TYR	33	35.645	99.125	34.999	1.00	15.00
	ATOM	251	CA	TYR	33	35.069	97.799	35.139	1.00	15.00
	ATOM	252	C	TYR	33	36.074	96.673	34.766	1.00	15.00
	ATOM	253	O	TYR	33	36.246	95.735	35.533	1.00	15.00
	ATOM	254	CB	TYR	33	33.833	97.736	34.299	1.00	15.00
60	ATOM	255	CG	TYR	33	32.825	98.811	34.662	1.00	20.00
	ATOM	256	CD1	TYR	33	32.813	100.049	34.037	1.00	20.00
	ATOM	257	CD2	TYR	33	31.860	98.556	35.628	1.00	20.00
	ATOM	258	CE1	TYR	33	31.870	101.004	34.361	1.00	20.00
	ATOM	259	CE2	TYR	33	30.911	99.504	35.958	1.00	20.00
65	ATOM	260	CZ	TYR	33	30.920	100.726	35.322	1.00	20.00
	ATOM	261	OH	TYR	33	29.976	101.674	35.647	1.00	20.00

500

	ATOM	262	N	CYS	34	36.731	96.775	33.598	1.00	15.00
	ATOM	263	CA	CYS	34	37.714	95.789	33.174	1.00	15.00
	ATOM	264	C	CYS	34	38.778	95.615	34.286	1.00	15.00
	ATOM	265	O	CYS	34	39.259	94.506	34.522	1.00	15.00
5	ATOM	266	CB	CYS	34	38.330	96.178	31.835	1.00	15.00
	ATOM	267	SG	CYS	34	37.180	96.140	30.425	1.00	20.00
	ATOM	268	N	GLN	35	39.116	96.718	34.948	1.00	15.00
	ATOM	269	CA	GLN	35	40.080	96.814	36.064	1.00	15.00
	ATOM	270	CB	GLN	35	40.330	98.299	36.335	1.00	15.00
10	ATOM	271	CG	GLN	35	41.572	98.628	37.123	1.00	15.00
	ATOM	272	CD	GLN	35	42.767	97.920	36.557	1.00	15.00
	ATOM	273	OE1	GLN	35	43.050	98.031	35.366	1.00	15.00
	ATOM	274	NE2	GLN	35	43.616	97.157	37.216	1.00	15.00
	ATOM	275	C	GLN	35	39.563	96.172	37.348	1.00	15.00
15	ATOM	276	O	GLN	35	40.278	95.410	38.005	1.00	15.00
	ATOM	277	N	ALA	36	38.312	96.490	37.661	1.00	15.00
	ATOM	278	CA	ALA	36	37.595	95.943	38.815	1.00	15.00
	ATOM	279	C	ALA	36	37.653	94.406	38.750	1.00	15.00
	ATOM	280	O	ALA	36	38.003	93.755	39.737	1.00	15.00
20	ATOM	281	CB	ALA	36	36.164	96.419	38.845	1.00	15.00
	ATOM	282	N	SER	37	37.275	93.858	37.569	1.00	15.00
	ATOM	283	CA	SER	37	37.258	92.382	37.440	1.00	15.00
	ATOM	284	C	SER	37	38.676	91.861	37.452	1.00	15.00
	ATOM	285	O	SER	37	38.962	90.810	38.051	1.00	15.00
25	ATOM	286	CB	SER	37	36.459	91.935	36.205	1.00	15.00
	ATOM	287	OG	SER	37	36.854	92.639	35.041	1.00	20.00
	ATOM	288	N	VAL	38	39.575	92.593	36.787	1.00	15.00
	ATOM	289	CA	VAL	38	40.964	92.230	36.763	1.00	15.00
	ATOM	290	C	VAL	38	41.528	92.053	38.178	1.00	15.00
30	ATOM	291	O	VAL	38	42.142	91.024	38.516	1.00	15.00
	ATOM	292	CB	VAL	38	41.797	93.318	36.023	1.00	15.00
	ATOM	293	CG1	VAL	38	43.296	93.111	36.239	1.00	20.00
	ATOM	294	CG2	VAL	38	41.483	93.300	34.530	1.00	20.00
	ATOM	295	N	THR	39	41.333	93.102	38.995	1.00	15.00
35	ATOM	296	CA	THR	39	41.833	93.097	40.368	1.00	15.00
	ATOM	297	C	THR	39	41.549	91.757	41.029	1.00	15.00
	ATOM	298	O	THR	39	41.979	91.511	42.160	1.00	15.00
	ATOM	299	CB	THR	39	41.161	94.224	41.168	1.00	15.00
	ATOM	300	OG1	THR	39	41.524	95.496	40.605	1.00	20.00
40	ATOM	301	CG2	THR	39	41.588	94.149	42.615	1.00	20.00
	END					-16.719	146.167	89.779	0.00	0.00

TABLE 18

	111							
	ATOM	1	C	CYS	1	11.122	61.996	7.481 1.00101.54
5	ATOM	2	O	CYS	1	9.901	61.868	7.536 1.00102.94
	ATOM	3	CB	CYS	1	12.164	63.950	6.316 1.00 96.91
	ATOM	4	SG	CYS	1	13.944	64.329	6.346 1.00 92.39
	ATOM	5	N	CYS	1	11.031	62.177	5.014 1.00 93.44
	ATOM	6	CA	CYS	1	11.828	62.447	6.217 1.00 98.53
10	ATOM	7	N	SER	2	11.973	61.781	8.490 1.00104.82
	ATOM	8	CA	SER	2	11.678	61.448	9.877 1.00107.58
	ATOM	9	CB	SER	2	10.647	60.316	10.041 1.00109.37
	ATOM	10	OG	SER	2	9.766	60.572	11.115 1.00110.15
	ATOM	11	C	SER	2	13.028	61.119	10.566 1.00108.57
15	ATOM	12	O	SER	2	13.853	62.024	10.707 1.00109.82
	ATOM	13	N	GLN	3	13.276	59.880	10.994 1.00108.52
	ATOM	14	CA	GLN	3	14.493	59.564	11.766 1.00109.56
	ATOM	15	CB	GLN	3	15.712	59.290	10.864 1.00108.23
	ATOM	16	CG	GLN	3	16.370	60.513	10.265 1.00109.80
20	ATOM	17	CD	GLN	3	15.996	60.607	8.813 1.00110.35
	ATOM	18	OE1	GLN	3	16.246	59.705	8.021 1.00110.92
	ATOM	19	NE2	GLN	3	15.381	61.632	8.247 1.00110.59
	ATOM	20	C	GLN	3	14.762	60.672	12.815 1.00110.79
	ATOM	21	O	GLN	3	15.839	61.273	12.861 1.00114.50
25	ATOM	22	N	ASN	4	13.744	60.918	13.656 1.00109.87
	ATOM	23	CA	ASN	4	13.675	61.874	14.793 1.00107.78
	ATOM	24	CB	ASN	4	14.393	61.290	16.002 1.00109.08
	ATOM	25	CG	ASN	4	15.801	60.785	15.707 1.00109.99
	ATOM	26	OD1	ASN	4	15.985	59.776	15.024 1.00109.73
30	ATOM	27	ND2	ASN	4	16.800	61.492	16.238 1.00109.43
	ATOM	28	C	ASN	4	14.160	63.290	14.466 1.00105.15
	ATOM	29	O	ASN	4	15.137	63.762	15.024 1.00103.42
	ATOM	30	N	GLU	5	13.465	63.971	13.520 1.00101.11
	ATOM	31	CA	GLU	5	13.837	65.365	13.103 1.00 95.31
35	ATOM	32	CB	GLU	5	14.976	65.332	12.083 1.00 92.47
	ATOM	33	CG	GLU	5	15.928	64.154	12.205 1.00 86.82
	ATOM	34	CD	GLU	5	16.169	63.547	10.837 1.00 85.55
	ATOM	35	OE1	GLU	5	15.190	63.396	10.086 1.00 85.72
	ATOM	36	OE2	GLU	5	17.331	63.236	10.538 1.00 81.01
40	ATOM	37	C	GLU	5	12.696	66.239	12.522 1.00 92.66
	ATOM	38	O	GLU	5	11.528	66.040	12.805 1.00 94.43
	ATOM	39	N	TYR	6	13.109	67.209	11.706 1.00 85.19
	ATOM	40	CA	TYR	6	12.200	68.163	11.067 1.00 78.14
	ATOM	41	CB	TYR	6	11.975	69.372	11.957 1.00 76.59
45	ATOM	42	CG	TYR	6	13.105	70.354	11.983 1.00 73.94
	ATOM	43	CD1	TYR	6	13.004	71.533	11.230 1.00 73.49
	ATOM	44	CE1	TYR	6	14.017	72.471	11.250 1.00 73.13
	ATOM	45	CD2	TYR	6	14.261	70.148	12.741 1.00 72.45
	ATOM	46	CE2	TYR	6	15.295	71.097	12.760 1.00 72.38
50	ATOM	47	CZ	TYR	6	15.159	72.248	12.022 1.00 74.15
	ATOM	48	OH	TYR	6	16.164	73.186	12.027 1.00 73.32
	ATOM	49	C	TYR	6	12.798	68.573	9.741 1.00 75.43
	ATOM	50	O	TYR	6	14.016	68.417	9.527 1.00 76.93
	ATOM	51	N	PHE	7	11.973	69.100	8.840 1.00 69.11
55	ATOM	52	CA	PHE	7	12.483	69.549	7.550 1.00 62.54
	ATOM	53	CB	PHE	7	11.548	69.070	6.435 1.00 61.14
	ATOM	54	CG	PHE	7	11.959	69.590	5.084 1.00 61.11
	ATOM	55	CD1	PHE	7	13.042	69.009	4.435 1.00 59.90
	ATOM	56	CD2	PHE	7	11.307	70.663	4.478 1.00 63.23
60	ATOM	57	CE1	PHE	7	13.472	69.491	3.198 1.00 61.45
	ATOM	58	CE2	PHE	7	11.725	71.161	3.238 1.00 61.15
	ATOM	59	CZ	PHE	7	12.808	70.576	2.596 1.00 61.89
	ATOM	60	C	PHE	7	12.644	71.077	7.517 1.00 60.79
	ATOM	61	O	PHE	7	11.678	71.797	7.250 1.00 63.10
65	ATOM	62	N	ASP	8	13.857	71.570	7.785 1.00 57.06
	ATOM	63	CA	ASP	8	14.128	73.013	7.779 1.00 52.20

	ATOM	64	CB	ASP	8	15.516	73.319	8.332	1.00	46.52
	ATOM	65	CG	ASP	8	15.776	74.793	8.611	1.00	47.00
	ATOM	66	OD1	ASP	8	15.241	75.630	7.852	1.00	49.00
	ATOM	67	OD2	ASP	8	16.514	75.095	9.571	1.00	48.43
5	ATOM	68	C	ASP	8	14.006	73.557	6.349	1.00	53.02
	ATOM	69	O	ASP	8	14.839	73.261	5.487	1.00	53.25
	ATOM	70	N	SER	9	12.960	74.347	6.102	1.00	55.34
	ATOM	71	CA	SER	9	12.730	74.941	4.779	1.00	54.13
	ATOM	72	CB	SER	9	11.366	75.637	4.724	1.00	54.65
10	ATOM	73	OG	SER	9	10.321	74.712	4.963	1.00	54.09
	ATOM	74	C	SER	9	13.813	75.955	4.404	1.00	54.96
	ATOM	75	O	SER	9	14.078	76.181	3.211	1.00	54.11
	ATOM	76	N	LEU	10	14.447	76.567	5.399	1.00	49.64
	ATOM	77	CA	LEU	10	15.498	77.514	5.088	1.00	46.11
15	ATOM	78	CB	LEU	10	16.011	78.213	6.339	1.00	40.39
	ATOM	79	CG	LEU	10	16.962	79.368	6.110	1.00	34.58
	ATOM	80	CD1	LEU	10	16.236	80.531	5.431	1.00	33.04
	ATOM	81	CD2	LEU	10	17.605	79.815	7.425	1.00	33.82
	ATOM	82	C	LEU	10	16.611	76.746	4.375	1.00	49.82
20	ATOM	83	O	LEU	10	17.124	77.190	3.361	1.00	50.94
	ATOM	84	N	LEU	11	16.984	75.588	4.920	1.00	51.20
	ATOM	85	CA	LEU	11	18.053	74.784	4.330	1.00	51.39
	ATOM	86	CB	LEU	11	18.766	73.971	5.417	1.00	47.82
	ATOM	87	CG	LEU	11	19.358	74.727	6.615	1.00	46.28
25	ATOM	88	CD1	LEU	11	20.070	73.757	7.551	1.00	49.44
	ATOM	89	CD2	LEU	11	20.315	75.819	6.159	1.00	44.53
	ATOM	90	C	LEU	11	17.607	73.840	3.208	1.00	53.87
	ATOM	91	O	LEU	11	18.411	73.434	2.379	1.00	50.27
	ATOM	92	N	HIS	12	16.319	73.478	3.192	1.00	58.57
30	ATOM	93	CA	HIS	12	15.775	72.557	2.215	1.00	64.33
	ATOM	94	CB	HIS	12	16.073	73.004	0.785	1.00	65.68
	ATOM	95	CG	HIS	12	15.349	74.319	0.396	1.00	69.20
	ATOM	96	CD2	HIS	12	14.164	74.836	0.804	1.00	69.34
	ATOM	97	ND1	HIS	12	15.885	75.239	-0.480	1.00	71.64
35	ATOM	98	CE1	HIS	12	15.064	76.273	-0.589	1.00	72.29
	ATOM	99	NE2	HIS	12	14.012	76.052	0.180	1.00	70.37
	ATOM	100	C	HIS	12	16.394	71.178	2.392	1.00	66.69
	ATOM	101	O	HIS	12	16.751	70.490	1.430	1.00	67.11
	ATOM	102	N	ALA	13	16.535	70.793	3.657	1.00	69.57
40	ATOM	103	CA	ALA	13	17.139	69.521	4.029	1.00	71.62
	ATOM	104	CB	ALA	13	18.643	69.636	4.114	1.00	69.66
	ATOM	105	C	ALA	13	16.555	69.077	5.367	1.00	73.99
	ATOM	106	O	ALA	13	15.668	69.744	5.912	1.00	75.45
	ATOM	107	N	CYS	14	17.056	67.969	5.908	1.00	75.75
45	ATOM	108	CA	CYS	14	16.534	67.462	7.171	1.00	77.01
	ATOM	109	C	CYS	14	17.491	67.571	8.339	1.00	75.86
	ATOM	110	O	CYS	14	18.587	67.002	8.345	1.00	73.08
	ATOM	111	CB	CYS	14	16.049	66.023	6.981	1.00	80.96
	ATOM	112	SG	CYS	14	14.259	65.862	6.705	1.00	86.58
50	ATOM	113	N	ILE	15	17.021	68.331	9.352	1.00	75.89
	ATOM	114	CA	ILE	15	17.820	68.581	10.541	1.00	77.02
	ATOM	115	CB	ILE	15	17.928	70.096	10.832	1.00	76.68
	ATOM	116	CG2	ILE	15	18.753	70.339	12.092	1.00	76.37
	ATOM	117	CG1	ILE	15	18.542	70.851	9.640	1.00	76.28
55	ATOM	118	CD1	ILE	15	19.984	70.495	9.345	1.00	74.77
	ATOM	119	C	ILE	15	17.298	67.852	11.800	1.00	78.19
	ATOM	120	O	ILE	15	16.115	67.832	12.108	1.00	77.54
	ATOM	121	N	PRO	16	18.297	67.270	12.517	1.00	80.09
	ATOM	122	CD	PRO	16	19.357	66.499	11.828	1.00	79.64
60	ATOM	123	CA	PRO	16	17.995	66.591	13.813	1.00	82.16
	ATOM	124	CB	PRO	16	19.363	66.094	14.245	1.00	80.00
	ATOM	125	CG	PRO	16	19.976	65.706	12.944	1.00	80.17
	ATOM	126	C	PRO	16	17.182	67.421	14.764	1.00	84.28
	ATOM	127	O	PRO	16	17.588	68.538	15.079	1.00	84.12
65	ATOM	128	N	CYS	17	16.032	66.929	15.237	1.00	87.37
	ATOM	129	CA	CYS	17	15.221	67.708	16.158	1.00	88.33

	ATOM	130	CB	CYS	17	14.085	66.857	16.718	1.00	86.90
	ATOM	131	SG	CYS	17	12.670	66.654	15.596	1.00	84.56
	ATOM	132	C	CYS	17	16.061	68.279	17.306	1.00	89.64
	ATOM	133	O	CYS	17	15.860	69.428	17.732	1.00	89.77
5	ATOM	134	N	GLN	18	16.994	67.464	17.800	1.00	91.08
	ATOM	135	CA	GLN	18	17.879	67.871	18.851	1.00	91.16
	ATOM	136	CB	GLN	18	19.146	67.016	18.899	1.00	92.68
	ATOM	137	CG	GLN	18	18.873	65.535	18.665	1.00	98.31
	ATOM	138	CD	GLN	18	20.085	64.733	18.174	1.00	101.37
10	ATOM	139	OE1	GLN	18	20.535	64.902	17.048	1.00	102.74
	ATOM	140	NE2	GLN	18	20.748	63.827	18.868	1.00	103.76
	ATOM	141	C	GLN	18	18.260	69.337	18.629	1.00	89.18
	ATOM	142	O	GLN	18	18.070	70.192	19.497	1.00	89.64
	ATOM	143	N	LEU	19	18.812	69.582	17.439	1.00	87.56
15	ATOM	144	CA	LEU	19	19.294	70.887	17.024	1.00	85.33
	ATOM	145	CB	LEU	19	19.648	70.861	15.565	1.00	79.98
	ATOM	146	CG	LEU	19	21.137	70.776	15.306	1.00	77.91
	ATOM	147	CD1	LEU	19	21.409	70.375	13.865	1.00	78.13
	ATOM	148	CD2	LEU	19	21.821	72.092	15.634	1.00	72.50
20	ATOM	149	C	LEU	19	18.344	72.042	17.295	1.00	86.05
	ATOM	150	O	LEU	19	18.774	73.096	17.789	1.00	88.06
	ATOM	151	N	ARG	20	17.079	71.871	16.998	1.00	85.11
	ATOM	152	CA	ARG	20	16.132	72.976	17.196	1.00	83.33
	ATOM	153	CB	ARG	20	14.926	72.805	16.276	1.00	78.93
25	ATOM	154	CG	ARG	20	14.925	73.718	15.053	1.00	73.68
	ATOM	155	CD	ARG	20	14.760	75.177	15.435	1.00	67.23
	ATOM	156	NE	ARG	20	13.502	75.733	14.935	1.00	63.84
	ATOM	157	CZ	ARG	20	13.079	75.616	13.674	1.00	64.39
	ATOM	158	NH1	ARG	20	13.826	74.958	12.793	1.00	66.99
30	ATOM	159	NH2	ARG	20	11.928	76.157	13.309	1.00	60.63
	ATOM	160	C	ARG	20	15.681	73.140	18.648	1.00	84.21
	ATOM	161	O	ARG	20	14.658	73.779	18.894	1.00	86.28
	ATOM	162	N	CYS	21	16.419	72.582	19.597	1.00	85.02
	ATOM	163	CA	CYS	21	16.024	72.758	20.990	1.00	83.51
35	ATOM	164	CB	CYS	21	15.917	71.386	21.670	1.00	81.05
	ATOM	165	SG	CYS	21	14.739	70.244	20.880	1.00	73.08
	ATOM	166	C	CYS	21	16.961	73.719	21.766	1.00	84.26
	ATOM	167	O	CYS	21	17.683	73.266	22.642	1.00	84.48
	ATOM	168	N	SER	22	16.973	75.060	21.437	1.00	84.61
40	ATOM	169	CA	SER	22	17.802	76.119	22.090	1.00	85.07
	ATOM	170	C	SER	22	16.915	77.082	22.866	1.00	85.65
	ATOM	171	O	SER	22	17.260	78.234	23.118	1.00	86.77
	ATOM	172	CB	SER	22	18.601	76.931	21.061	1.00	20.00
	ATOM	173	OG	SER	22	17.832	78.008	20.562	1.00	20.00
45	ATOM	174	N	SER	23	15.784	76.538	23.223	1.00	84.19
	ATOM	175	CA	SER	23	14.816	77.255	24.023	1.00	82.44
	ATOM	176	C	SER	23	13.644	77.820	23.214	1.00	84.32
	ATOM	177	O	SER	23	12.711	77.071	22.921	1.00	81.24
	ATOM	178	CB	SER	23	15.555	78.331	24.819	1.00	78.00
50	ATOM	179	OG	SER	23	14.777	79.504	24.950	1.00	20.00
	ATOM	180	N	ASN	24	13.648	79.072	22.851	1.00	87.06
	ATOM	181	CA	ASN	24	12.491	79.490	22.086	1.00	88.17
	ATOM	182	C	ASN	24	12.444	78.687	20.777	1.00	88.45
	ATOM	183	O	ASN	24	11.599	78.949	19.940	1.00	84.89
55	ATOM	184	CB	ASN	24	12.565	80.949	21.695	0.00	86.84
	ATOM	185	CG	ASN	24	12.788	81.844	22.860	0.00	85.55
	ATOM	186	OD1	ASN	24	12.409	81.508	23.979	1.00	20.00
	ATOM	187	ND2	ASN	24	13.399	82.998	22.629	1.00	20.00
	ATOM	188	N	THR	25	13.371	77.706	20.611	1.00	89.49
60	ATOM	189	CA	THR	25	13.456	77.034	19.297	1.00	91.86
	ATOM	190	C	THR	25	12.877	75.593	19.091	1.00	95.75
	ATOM	191	O	THR	25	13.333	74.835	18.218	1.00	96.59
	ATOM	192	CB	THR	25	14.893	77.129	18.834	0.00	91.23
	ATOM	193	OG1	THR	25	15.293	78.501	18.828	1.00	20.00
65	ATOM	194	CG2	THR	25	15.032	76.535	17.449	1.00	20.00
	ATOM	195	N	PRO	26	11.870	75.224	19.916	1.00	98.41

	ATOM	196	CA	PRO	26	11.022	73.922	19.739	1.00	96.39
	ATOM	197	C	PRO	26	9.888	73.735	18.538	1.00	96.88
	ATOM	198	O	PRO	26	8.751	74.141	18.792	1.00	94.52
	ATOM	199	CB	PRO	26	10.409	73.811	21.122	0.00	95.29
5	ATOM	200	CG	PRO	26	11.493	74.273	22.049	0.00	92.34
	ATOM	201	CD	PRO	26	12.408	75.168	21.274	0.00	89.50
	ATOM	202	N	PRO	27	10.157	73.172	17.255	1.00	97.72
	ATOM	203	CA	PRO	27	9.094	72.880	16.132	1.00	98.94
	ATOM	204	C	PRO	27	7.876	71.877	16.324	1.00	100.71
10	ATOM	205	O	PRO	27	7.909	71.102	17.277	1.00	100.56
	ATOM	206	CB	PRO	27	9.979	72.563	14.948	0.00	97.33
	ATOM	207	CG	PRO	27	11.113	73.518	15.130	0.00	94.51
	ATOM	208	CD	PRO	27	11.238	73.857	16.571	0.00	92.58
	ATOM	209	N	LEU	28	6.781	71.846	15.465	1.00	15.00
15	ATOM	210	CA	LEU	28	5.805	70.727	15.681	1.00	15.00
	ATOM	211	C	LEU	28	6.773	69.547	15.672	1.00	15.00
	ATOM	212	O	LEU	28	7.972	69.735	15.884	1.00	15.00
	ATOM	213	CB	LEU	28	4.715	70.597	14.563	1.00	15.00
	ATOM	214	CG	LEU	28	3.215	70.751	14.968	1.00	15.00
20	ATOM	215	CD1	LEU	28	2.492	69.436	14.799	1.00	15.00
	ATOM	216	CD2	LEU	28	3.094	71.262	16.403	1.00	15.00
	ATOM	217	N	THR	29	6.258	68.361	15.395	1.00	15.00
	ATOM	218	CA	THR	29	7.139	67.205	15.277	1.00	15.00
	ATOM	219	C	THR	29	8.448	67.451	16.020	1.00	15.00
25	ATOM	220	O	THR	29	9.461	66.840	15.721	1.00	15.00
	ATOM	221	CB	THR	29	7.391	66.889	13.798	1.00	15.00
	ATOM	222	OG1	THR	29	6.548	65.810	13.404	1.00	20.00
	ATOM	223	CG2	THR	29	8.853	66.518	13.589	1.00	20.00
	ATOM	224	N	CYS	30	8.373	68.362	17.006	1.00	15.00
30	ATOM	225	CA	CYS	30	9.571	68.704	17.746	1.00	15.00
	ATOM	226	C	CYS	30	9.258	68.868	19.228	1.00	15.00
	ATOM	227	O	CYS	30	10.111	68.615	20.062	1.00	15.00
	ATOM	228	CB	CYS	30	10.244	69.924	17.135	1.00	15.00
	ATOM	229	SG	CYS	30	11.088	69.622	15.551	1.00	20.00
35	ATOM	230	N	GLN	31	8.029	69.283	19.535	1.00	15.00
	ATOM	231	CA	GLN	31	7.645	69.418	20.920	1.00	15.00
	ATOM	232	C	GLN	31	7.786	68.091	21.669	1.00	15.00
	ATOM	233	O	GLN	31	8.385	68.016	22.746	1.00	15.00
	ATOM	234	CB	GLN	31	6.237	69.945	21.039	1.00	15.00
40	ATOM	235	CG	GLN	31	6.214	71.467	20.991	1.00	20.00
	ATOM	236	CD	GLN	31	4.829	72.040	21.213	1.00	20.00
	ATOM	237	OE1	GLN	31	3.944	71.365	21.738	1.00	20.00
	ATOM	238	NE2	GLN	31	4.421	73.263	20.891	1.00	20.00
	ATOM	239	N	ARG	32	7.242	67.055	21.094	1.00	15.00
45	ATOM	240	CA	ARG	32	7.325	65.718	21.644	1.00	15.00
	ATOM	241	C	ARG	32	8.771	65.271	21.835	1.00	15.00
	ATOM	242	O	ARG	32	9.121	64.639	22.847	1.00	15.00
	ATOM	243	CB	ARG	32	6.519	64.741	20.762	1.00	15.00
	ATOM	244	CG	ARG	32	5.021	65.051	20.664	1.00	15.00
50	ATOM	245	CD	ARG	32	4.309	63.984	19.839	1.00	15.00
	ATOM	246	NE	ARG	32	2.892	64.268	19.651	1.00	20.00
	ATOM	247	CZ	ARG	32	2.048	63.459	19.020	1.00	20.00
	ATOM	248	NH1	ARG	32	2.485	62.311	18.517	1.00	20.00
	ATOM	249	NH2	ARG	32	0.772	63.794	18.895	1.00	20.00
55	ATOM	250	N	TYR	33	9.593	65.632	20.852	1.00	15.00
	ATOM	251	CA	TYR	33	11.007	65.302	20.889	1.00	15.00
	ATOM	252	C	TYR	33	11.757	66.033	22.038	1.00	15.00
	ATOM	253	O	TYR	33	12.473	65.394	22.797	1.00	15.00
	ATOM	254	CB	TYR	33	11.596	65.623	19.553	1.00	15.00
60	ATOM	255	CG	TYR	33	10.896	64.901	18.417	1.00	20.00
	ATOM	256	CD1	TYR	33	9.827	65.464	17.735	1.00	20.00
	ATOM	257	CD2	TYR	33	11.342	63.648	18.014	1.00	20.00
	ATOM	258	CE1	TYR	33	9.221	64.803	16.685	1.00	20.00
	ATOM	259	CE2	TYR	33	10.744	62.980	16.964	1.00	20.00
65	ATOM	260	CZ	TYR	33	9.684	63.561	16.303	1.00	20.00
	ATOM	261	OH	TYR	33	9.085	62.899	15.255	1.00	20.00

	ATOM	262	N	CYS	34	11.584	67.360	22.160	1.00	15.00
	ATOM	263	CA	CYS	34	12.218	68.129	23.221	1.00	15.00
	ATOM	264	C	CYS	34	11.874	67.494	24.590	1.00	15.00
	ATOM	265	O	CYS	34	12.711	67.465	25.493	1.00	15.00
5	ATOM	266	CB	CYS	34	11.814	69.597	23.142	1.00	15.00
	ATOM	267	SG	CYS	34	12.423	70.480	21.672	1.00	20.00
	ATOM	268	N	GLN	35	10.647	66.993	24.709	1.00	15.00
	ATOM	269	CA	GLN	35	10.086	66.313	25.896	1.00	15.00
	ATOM	270	CB	GLN	35	8.583	66.139	25.663	1.00	15.00
10	ATOM	271	CG	GLN	35	7.751	65.865	26.889	1.00	15.00
	ATOM	272	CD	GLN	35	8.071	66.842	27.981	1.00	15.00
	ATOM	273	OE1	GLN	35	8.022	68.051	27.769	1.00	15.00
	ATOM	274	NE2	GLN	35	8.406	66.553	29.223	1.00	15.00
	ATOM	275	C	GLN	35	10.698	64.936	26.133	1.00	15.00
15	ATOM	276	O	GLN	35	11.078	64.600	27.258	1.00	15.00
	ATOM	277	N	ALA	36	10.792	64.178	25.046	1.00	15.00
	ATOM	278	CA	ALA	36	11.401	62.846	25.031	1.00	15.00
	ATOM	279	C	ALA	36	12.819	62.944	25.620	1.00	15.00
	ATOM	280	O	ALA	36	13.180	62.164	26.505	1.00	15.00
20	ATOM	281	CB	ALA	36	11.445	62.282	23.632	1.00	15.00
	ATOM	282	N	SER	37	13.609	63.906	25.082	1.00	15.00
	ATOM	283	CA	SER	37	15.005	64.035	25.563	1.00	15.00
	ATOM	284	C	SER	37	15.001	64.555	26.981	1.00	15.00
	ATOM	285	O	SER	37	15.805	64.116	27.821	1.00	15.00
25	ATOM	286	CB	SER	37	15.851	64.889	24.606	1.00	15.00
	ATOM	287	OG	SER	37	15.208	66.109	24.283	1.00	20.00
	ATOM	288	N	VAL	38	14.095	65.499	27.257	1.00	15.00
	ATOM	289	CA	VAL	38	13.959	66.041	28.579	1.00	15.00
	ATOM	290	C	VAL	38	13.751	64.938	29.625	1.00	15.00
30	ATOM	291	O	VAL	38	14.455	64.863	30.649	1.00	15.00
	ATOM	292	CB	VAL	38	12.755	67.026	28.642	1.00	15.00
	ATOM	293	CG1	VAL	38	12.405	67.384	30.086	1.00	20.00
	ATOM	294	CG2	VAL	38	13.069	68.295	27.856	1.00	20.00
	ATOM	295	N	THR	39	12.738	64.096	29.358	1.00	15.00
35	ATOM	296	CA	THR	39	12.397	63.007	30.270	1.00	15.00
	ATOM	297	C	THR	39	13.658	62.302	30.746	1.00	15.00
	ATOM	298	O	THR	39	13.596	61.414	31.602	1.00	15.00
	ATOM	299	CB	THR	39	11.478	62.003	29.557	1.00	15.00
	ATOM	300	OG1	THR	39	10.241	62.647	29.209	1.00	20.00
40	ATOM	301	CG2	THR	39	11.217	60.819	30.458	1.00	20.00
	END					-23.152	-5.163	-22.411	0.00	0.00

TABLE 19

	111							
	ATOM	1	C	CYS	1	-49.816	59.772	-22.893 1.00101.54
5	ATOM	2	O	CYS	1	-49.277	59.078	-23.750 1.00102.94
	ATOM	3	CB	CYS	1	-49.505	62.255	-22.858 1.00 96.91
	ATOM	4	SG	CYS	1	-50.410	63.385	-21.754 1.00 92.39
	ATOM	5	N	CYS	1	-47.634	60.753	-22.257 1.00 93.44
	ATOM	6	CA	CYS	1	-49.092	60.918	-22.210 1.00 98.53
10	ATOM	7	N	SER	2	-51.075	59.639	-22.462 1.00104.82
	ATOM	8	CA	SER	2	-52.113	58.739	-22.945 1.00107.58
	ATOM	9	CB	SER	2	-51.646	57.282	-23.117 1.00109.37
	ATOM	10	OG	SER	2	-52.201	56.695	-24.276 1.00110.15
	ATOM	11	C	SER	2	-53.314	58.860	-21.971 1.00108.57
15	ATOM	12	O	SER	2	-53.927	59.929	-21.922 1.00109.82
	ATOM	13	N	GLN	3	-53.667	57.822	-21.210 1.00108.52
	ATOM	14	CA	GLN	3	-54.879	57.862	-20.370 1.00109.56
	ATOM	15	CB	GLN	3	-54.631	58.531	-19.005 1.00108.23
	ATOM	16	CG	GLN	3	-54.553	60.043	-19.022 1.00109.80
20	ATOM	17	CD	GLN	3	-53.117	60.458	-18.880 1.00110.35
	ATOM	18	OE1	GLN	3	-52.439	60.127	-17.914 1.00110.92
	ATOM	19	NE2	GLN	3	-52.447	61.201	-19.747 1.00110.59
	ATOM	20	C	GLN	3	-56.048	58.512	-21.152 1.00110.79
	ATOM	21	O	GLN	3	-56.662	59.487	-20.708 1.00114.50
25	ATOM	22	N	ASN	4	-56.334	57.936	-22.332 1.00109.87
	ATOM	23	CA	ASN	4	-57.404	58.273	-23.307 1.00107.78
	ATOM	24	CB	ASN	4	-58.734	57.708	-22.829 1.00109.08
	ATOM	25	CG	ASN	4	-59.081	58.063	-21.387 1.00109.99
	ATOM	26	OD1	ASN	4	-58.456	57.576	-20.444 1.00109.73
30	ATOM	27	ND2	ASN	4	-60.095	58.915	-21.222 1.00109.43
	ATOM	28	C	ASN	4	-57.504	59.762	-23.654 1.00105.15
	ATOM	29	O	ASN	4	-58.505	60.404	-23.378 1.00103.42
	ATOM	30	N	GLU	5	-56.426	60.324	-24.257 1.00101.11
	ATOM	31	CA	GLU	5	-56.392	61.774	-24.646 1.00 95.31
35	ATOM	32	CB	GLU	5	-56.030	62.643	-23.441 1.00 92.47
	ATOM	33	CG	GLU	5	-56.453	62.092	-22.088 1.00 86.82
	ATOM	34	CD	GLU	5	-55.300	62.199	-21.110 1.00 85.55
	ATOM	35	OE1	GLU	5	-54.166	61.885	-21.509 1.00 85.72
	ATOM	36	OE2	GLU	5	-55.549	62.598	-19.963 1.00 81.01
40	ATOM	37	C	GLU	5	-55.446	62.152	-25.813 1.00 92.66
	ATOM	38	O	GLU	5	-55.122	61.344	-26.666 1.00 94.43
	ATOM	39	N	TYR	6	-55.035	63.420	-25.797 1.00 85.19
	ATOM	40	CA	TYR	6	-54.155	63.994	-26.819 1.00 78.14
	ATOM	41	CB	TYR	6	-54.964	64.550	-27.975 1.00 76.59
45	ATOM	42	CG	TYR	6	-55.628	65.865	-27.706 1.00 73.94
	ATOM	43	CD1	TYR	6	-55.054	67.039	-28.215 1.00 73.49
	ATOM	44	CE1	TYR	6	-55.652	68.265	-28.004 1.00 73.13
	ATOM	45	CD2	TYR	6	-56.811	65.971	-26.970 1.00 72.45
	ATOM	46	CE2	TYR	6	-57.421	67.215	-26.750 1.00 72.38
50	ATOM	47	CZ	TYR	6	-56.840	68.345	-27.273 1.00 74.15
	ATOM	48	OH	TYR	6	-57.422	69.573	-27.065 1.00 73.32
	ATOM	49	C	TYR	6	-53.321	65.076	-26.172 1.00 75.43
	ATOM	50	O	TYR	6	-53.688	65.596	-25.102 1.00 76.93
	ATOM	51	N	PHE	7	-52.204	65.436	-26.799 1.00 69.11
55	ATOM	52	CA	PHE	7	-51.364	66.496	-26.254 1.00 62.54
	ATOM	53	CB	PHE	7	-49.894	66.067	-26.312 1.00 61.14
	ATOM	54	CG	PHE	7	-48.964	67.160	-25.862 1.00 61.11
	ATOM	55	CD1	PHE	7	-48.839	67.427	-24.504 1.00 59.90
	ATOM	56	CD2	PHE	7	-48.247	67.938	-26.770 1.00 63.23
60	ATOM	57	CE1	PHE	7	-48.012	68.458	-24.055 1.00 61.45
	ATOM	58	CE2	PHE	7	-47.413	68.977	-26.338 1.00 61.15
	ATOM	59	CZ	PHE	7	-47.295	69.238	-24.981 1.00 61.89
	ATOM	60	C	PHE	7	-51.582	67.820	-27.004 1.00 60.79
	ATOM	61	O	PHE	7	-50.976	68.044	-28.055 1.00 63.10
65	ATOM	62	N	ASP	8	-52.442	68.692	-26.469 1.00 57.06
	ATOM	63	CA	ASP	8	-52.727	69.988	-27.097 1.00 52.20

	ATOM	64	CB	ASP	8	-53.898	70.689	-26.416	1.00	46.52
	ATOM	65	CG	ASP	8	-54.429	71.904	-27.163	1.00	47.00
	ATOM	66	OD1	ASP	8	-53.607	72.601	-27.798	1.00	49.00
	ATOM	67	OD2	ASP	8	-55.650	72.154	-27.099	1.00	48.43
5	ATOM	68	C	ASP	8	-51.478	70.879	-27.034	1.00	53.02
	ATOM	69	O	ASP	8	-51.081	71.336	-25.957	1.00	53.25
	ATOM	70	N	SER	9	-50.859	71.116	-28.191	1.00	55.34
	ATOM	71	CA	SER	9	-49.659	71.958	-28.272	1.00	54.13
	ATOM	72	CB	SER	9	-49.048	71.903	-29.676	1.00	54.65
10	ATOM	73	OG	SER	9	-48.663	70.580	-30.003	1.00	54.09
	ATOM	74	C	SER	9	-49.953	73.419	-27.924	1.00	54.96
	ATOM	75	O	SER	9	-49.059	74.150	-27.465	1.00	54.11
	ATOM	76	N	LEU	10	-51.192	73.858	-28.128	1.00	49.64
	ATOM	77	CA	LEU	10	-51.520	75.226	-27.787	1.00	46.11
15	ATOM	78	CB	LEU	10	-52.934	75.589	-28.216	1.00	40.39
	ATOM	79	CG	LEU	10	-53.311	77.049	-28.095	1.00	34.58
	ATOM	80	CD1	LEU	10	-52.505	77.891	-29.087	1.00	33.04
	ATOM	81	CD2	LEU	10	-54.814	77.245	-28.303	1.00	33.82
	ATOM	82	C	LEU	10	-51.332	75.380	-26.277	1.00	49.82
20	ATOM	83	O	LEU	10	-50.735	76.339	-25.817	1.00	50.94
	ATOM	84	N	LEU	11	-51.854	74.424	-25.508	1.00	51.20
	ATOM	85	CA	LEU	11	-51.750	74.484	-24.052	1.00	51.39
	ATOM	86	CB	LEU	11	-52.945	73.774	-23.405	1.00	47.82
	ATOM	87	CG	LEU	11	-54.357	74.237	-23.791	1.00	46.28
25	ATOM	88	CD1	LEU	11	-55.402	73.454	-23.005	1.00	49.44
	ATOM	89	CD2	LEU	11	-54.530	75.731	-23.555	1.00	44.53
	ATOM	90	C	LEU	11	-50.452	73.909	-23.473	1.00	53.87
	ATOM	91	O	LEU	11	-50.058	74.252	-22.366	1.00	50.27
	ATOM	92	N	HIS	12	-49.792	73.019	-24.223	1.00	58.57
30	ATOM	93	CA	HIS	12	-48.578	72.366	-23.779	1.00	64.33
	ATOM	94	CB	HIS	12	-47.516	73.375	-23.345	1.00	65.68
	ATOM	95	CG	HIS	12	-46.982	74.238	-24.518	1.00	69.20
	ATOM	96	CD2	HIS	12	-46.841	73.956	-25.836	1.00	69.34
	ATOM	97	ND1	HIS	12	-46.571	75.545	-24.361	1.00	71.64
35	ATOM	98	CE1	HIS	12	-46.205	76.035	-25.536	1.00	72.29
	ATOM	99	NE2	HIS	12	-46.360	75.091	-26.448	1.00	70.37
	ATOM	100	C	HIS	12	-48.869	71.477	-22.578	1.00	66.69
	ATOM	101	O	HIS	12	-48.117	71.429	-21.599	1.00	67.11
	ATOM	102	N	ALA	13	-49.999	70.783	-22.665	1.00	69.57
40	ATOM	103	CA	ALA	13	-50.466	69.905	-21.601	1.00	71.62
	ATOM	104	CB	ALA	13	-51.260	70.672	-20.569	1.00	69.66
	ATOM	105	C	ALA	13	-51.313	68.796	-22.220	1.00	73.99
	ATOM	106	O	ALA	13	-51.449	68.727	-23.447	1.00	75.45
	ATOM	107	N	CYS	14	-51.898	67.943	-21.382	1.00	75.75
45	ATOM	108	CA	CYS	14	-52.702	66.839	-21.894	1.00	77.01
	ATOM	109	C	CYS	14	-54.186	66.961	-21.624	1.00	75.86
	ATOM	110	O	CYS	14	-54.643	67.012	-20.478	1.00	73.08
	ATOM	111	CB	CYS	14	-52.146	65.516	-21.365	1.00	80.96
	ATOM	112	SG	CYS	14	-51.044	64.645	-22.521	1.00	86.58
50	ATOM	113	N	ILE	15	-54.937	66.997	-22.746	1.00	75.89
	ATOM	114	CA	ILE	15	-56.382	67.151	-22.682	1.00	77.02
	ATOM	115	CB	ILE	15	-56.858	68.324	-23.571	1.00	76.68
	ATOM	116	CG2	ILE	15	-58.376	68.459	-23.506	1.00	76.37
	ATOM	117	CG1	ILE	15	-56.187	69.646	-23.156	1.00	76.28
55	ATOM	118	CD1	ILE	15	-56.566	70.137	-21.775	1.00	74.77
	ATOM	119	C	ILE	15	-57.157	65.869	-23.064	1.00	78.19
	ATOM	120	O	ILE	15	-56.869	65.189	-24.039	1.00	77.54
	ATOM	121	N	PRO	16	-58.189	65.611	-22.216	1.00	80.09
	ATOM	122	CD	PRO	16	-57.997	65.729	-20.752	1.00	79.64
60	ATOM	123	CA	PRO	16	-59.106	64.460	-22.475	1.00	82.16
	ATOM	124	CB	PRO	16	-60.071	64.546	-21.306	1.00	80.00
	ATOM	125	CG	PRO	16	-59.176	64.981	-20.197	1.00	80.17
	ATOM	126	C	PRO	16	-59.649	64.414	-23.873	1.00	84.28
	ATOM	127	O	PRO	16	-60.241	65.396	-24.317	1.00	84.12
65	ATOM	128	N	CYS	17	-59.468	63.309	-24.605	1.00	87.37
	ATOM	129	CA	CYS	17	-59.981	63.232	-25.963	1.00	88.33

	ATOM	130	CB	CYS	17	-59.842	61.813	-26.505	1.00	86.90
	ATOM	131	SG	CYS	17	-58.171	61.384	-27.080	1.00	84.56
	ATOM	132	C	CYS	17	-61.445	63.680	-26.040	1.00	89.64
	ATOM	133	O	CYS	17	-61.853	64.365	-26.993	1.00	89.77
5	ATOM	134	N	GLN	18	-62.224	63.282	-25.034	1.00	91.08
	ATOM	135	CA	GLN	18	-63.606	63.653	-24.950	1.00	91.16
	ATOM	136	CB	GLN	18	-64.148	63.538	-23.524	1.00	92.68
	ATOM	137	CG	GLN	18	-63.648	62.294	-22.798	1.00	98.31
	ATOM	138	CD	GLN	18	-63.698	62.388	-21.268	1.00	101.37
10	ATOM	139	OE1	GLN	18	-62.942	63.136	-20.659	1.00	102.74
	ATOM	140	NE2	GLN	18	-64.514	61.719	-20.476	1.00	103.76
	ATOM	141	C	GLN	18	-63.756	65.096	-25.438	1.00	89.18
	ATOM	142	O	GLN	18	-64.522	65.391	-26.358	1.00	89.64
	ATOM	143	N	LEU	19	-63.000	65.975	-24.777	1.00	87.56
15	ATOM	144	CA	LEU	19	-63.010	67.405	-25.033	1.00	85.33
	ATOM	145	CB	LEU	19	-61.896	68.068	-24.276	1.00	79.98
	ATOM	146	CG	LEU	19	-62.359	68.788	-23.027	1.00	77.91
	ATOM	147	CD1	LEU	19	-61.180	69.102	-22.120	1.00	78.13
	ATOM	148	CD2	LEU	19	-63.115	70.056	-23.382	1.00	72.50
20	ATOM	149	C	LEU	19	-62.930	67.798	-26.499	1.00	86.05
	ATOM	150	O	LEU	19	-63.683	68.677	-26.947	1.00	88.06
	ATOM	151	N	ARG	20	-62.056	67.173	-27.250	1.00	85.11
	ATOM	152	CA	ARG	20	-61.909	67.552	-28.660	1.00	83.33
	ATOM	153	CB	ARG	20	-60.517	67.176	-29.163	1.00	78.93
25	ATOM	154	CG	ARG	20	-59.549	68.349	-29.289	1.00	73.68
	ATOM	155	CD	ARG	20	-59.970	69.316	-30.380	1.00	67.23
	ATOM	156	NE	ARG	20	-59.003	69.354	-31.478	1.00	63.84
	ATOM	157	CZ	ARG	20	-57.687	69.509	-31.314	1.00	64.39
	ATOM	158	NH1	ARG	20	-57.193	69.639	-30.087	1.00	66.99
30	ATOM	159	NH2	ARG	20	-56.888	69.538	-32.368	1.00	60.63
	ATOM	160	C	ARG	20	-62.987	66.959	-29.569	1.00	84.21
	ATOM	161	O	ARG	20	-62.793	66.910	-30.783	1.00	86.28
	ATOM	162	N	CYS	21	-64.102	66.516	-29.005	1.00	85.02
	ATOM	163	CA	CYS	21	-65.156	65.979	-29.859	1.00	83.51
35	ATOM	164	CB	CYS	21	-65.547	64.577	-29.369	1.00	81.05
	ATOM	165	SG	CYS	21	-64.173	63.382	-29.333	1.00	73.08
	ATOM	166	C	CYS	21	-66.383	66.919	-29.966	1.00	84.26
	ATOM	167	O	CYS	21	-67.439	66.580	-29.451	1.00	84.48
	ATOM	168	N	SER	22	-66.252	68.126	-30.623	1.00	84.61
40	ATOM	169	CA	SER	22	-67.332	69.140	-30.827	1.00	85.07
	ATOM	170	C	SER	22	-67.702	69.228	-32.301	1.00	85.65
	ATOM	171	O	SER	22	-68.214	70.232	-32.790	1.00	86.77
	ATOM	172	CB	SER	22	-66.897	70.536	-30.361	1.00	20.00
	ATOM	173	OG	SER	22	-66.220	71.225	-31.394	1.00	20.00
45	ATOM	174	N	SER	23	-67.422	68.132	-32.951	1.00	84.19
	ATOM	175	CA	SER	23	-67.749	67.974	-34.350	1.00	82.44
	ATOM	176	C	SER	23	-66.553	68.170	-35.289	1.00	84.32
	ATOM	177	O	SER	23	-65.774	67.230	-35.458	1.00	81.24
	ATOM	178	CB	SER	23	-68.914	68.908	-34.679	1.00	78.00
50	ATOM	179	OG	SER	23	-68.795	69.446	-35.981	1.00	20.00
	ATOM	180	N	ASN	24	-66.378	69.313	-35.889	1.00	87.06
	ATOM	181	CA	ASN	24	-65.211	69.381	-36.745	1.00	88.17
	ATOM	182	C	ASN	24	-63.953	69.174	-35.888	1.00	88.45
	ATOM	183	O	ASN	24	-62.852	69.288	-36.397	1.00	84.89
55	ATOM	184	CB	ASN	24	-65.067	70.736	-37.403	0.00	86.84
	ATOM	185	CG	ASN	24	-66.293	71.150	-38.134	0.00	85.55
	ATOM	186	OD1	ASN	24	-67.057	70.303	-38.591	1.00	20.00
	ATOM	187	ND2	ASN	24	-66.507	72.452	-38.267	1.00	20.00
	ATOM	188	N	THR	25	-64.132	68.874	-34.574	1.00	89.49
60	ATOM	189	CA	THR	25	-62.947	68.836	-33.692	1.00	91.86
	ATOM	190	C	THR	25	-62.332	67.472	-33.227	1.00	95.75
	ATOM	191	O	THR	25	-61.697	67.383	-32.163	1.00	96.59
	ATOM	192	CB	THR	25	-63.227	69.751	-32.520	0.00	91.23
	ATOM	193	OG1	THR	25	-63.564	71.050	-33.010	1.00	20.00
65	ATOM	194	CG2	THR	25	-62.013	69.827	-31.620	1.00	20.00
	ATOM	195	N	PRO	26	-62.540	66.410	-34.038	1.00	98.41

	ATOM	196	CA	PRO	26	-61.841	65.022	-33.863	1.00	96.39
	ATOM	197	C	PRO	26	-60.235	64.766	-34.210	1.00	96.88
	ATOM	198	O	PRO	26	-59.969	64.471	-35.378	1.00	94.52
	ATOM	199	CB	PRO	26	-62.751	64.154	-34.711	0.00	95.29
5	ATOM	200	CG	PRO	26	-64.124	64.707	-34.470	0.00	92.34
	ATOM	201	CD	PRO	26	-63.976	66.134	-34.048	0.00	89.50
	ATOM	202	N	PRO	27	-59.175	64.892	-33.263	1.00	97.72
	ATOM	203	CA	PRO	27	-57.660	64.556	-33.523	1.00	98.94
	ATOM	204	C	PRO	27	-57.143	63.107	-33.919	1.00100	71
10	ATOM	205	O	PRO	27	-57.906	62.157	-33.759	1.00100	56
	ATOM	206	CB	PRO	27	-57.003	65.135	-32.290	0.00	97.33
	ATOM	207	CG	PRO	27	-57.803	66.375	-32.052	0.00	94.51
	ATOM	208	CD	PRO	27	-59.161	66.195	-32.626	0.00	92.58
	ATOM	209	N	LEU	28	-55.873	62.874	-34.438	1.00	15.00
15	ATOM	210	CA	LEU	28	-55.478	61.435	-34.594	1.00	15.00
	ATOM	211	C	LEU	28	-55.792	60.936	-33.185	1.00	15.00
	ATOM	212	O	LEU	28	-56.562	61.575	-32.467	1.00	15.00
	ATOM	213	CB	LEU	28	-53.972	61.216	-34.968	1.00	15.00
	ATOM	214	CG	LEU	28	-53.639	60.494	-36.312	1.00	15.00
20	ATOM	215	CD1	LEU	28	-53.004	59.151	-36.039	1.00	15.00
	ATOM	216	CD2	LEU	28	-54.896	60.341	-37.168	1.00	15.00
	ATOM	217	N	THR	29	-55.175	59.834	-32.796	1.00	15.00
	ATOM	218	CA	THR	29	-55.356	59.353	-31.431	1.00	15.00
	ATOM	219	C	THR	29	-56.649	59.901	-30.838	1.00	15.00
25	ATOM	220	O	THR	29	-56.794	59.988	-29.629	1.00	15.00
	ATOM	221	CB	THR	29	-54.143	59.741	-30.576	1.00	15.00
	ATOM	222	OG1	THR	29	-53.281	58.613	-30.456	1.00	20.00
	ATOM	223	CG2	THR	29	-54.606	60.200	-29.200	1.00	20.00
	ATOM	224	N	CYS	30	-57.578	60.253	-31.742	1.00	15.00
30	ATOM	225	CA	CYS	30	-58.828	60.828	-31.285	1.00	15.00
	ATOM	226	C	CYS	30	-59.997	60.289	-32.100	1.00	15.00
	ATOM	227	O	CYS	30	-61.100	60.186	-31.588	1.00	15.00
	ATOM	228	CB	CYS	30	-58.746	62.348	-31.286	1.00	15.00
	ATOM	229	SG	CYS	30	-57.722	63.061	-29.960	1.00	20.00
35	ATOM	230	N	GLN	31	-59.735	59.939	-33.359	1.00	15.00
	ATOM	231	CA	GLN	31	-60.782	59.377	-34.178	1.00	15.00
	ATOM	232	C	GLN	31	-61.355	58.103	-33.552	1.00	15.00
	ATOM	233	O	GLN	31	-62.572	57.940	-33.414	1.00	15.00
	ATOM	234	CB	GLN	31	-60.284	59.102	-35.574	1.00	15.00
40	ATOM	235	CG	GLN	31	-60.402	60.339	-36.454	1.00	20.00
	ATOM	236	CD	GLN	31	-60.010	60.075	-37.894	1.00	20.00
	ATOM	237	OE1	GLN	31	-59.978	58.928	-38.339	1.00	20.00
	ATOM	238	NE2	GLN	31	-59.673	60.988	-38.799	1.00	20.00
	ATOM	239	N	ARG	32	-60.480	57.215	-33.173	1.00	15.00
45	ATOM	240	CA	ARG	32	-60.851	55.976	-32.520	1.00	15.00
	ATOM	241	C	ARG	32	-61.647	56.224	-31.243	1.00	15.00
	ATOM	242	O	ARG	32	-62.626	55.517	-30.948	1.00	15.00
	ATOM	243	CB	ARG	32	-59.590	55.122	-32.271	1.00	15.00
	ATOM	244	CG	ARG	32	-58.835	54.706	-33.538	1.00	15.00
50	ATOM	245	CD	ARG	32	-57.658	53.802	-33.185	1.00	15.00
	ATOM	246	NE	ARG	32	-56.859	53.435	-34.348	1.00	20.00
	ATOM	247	CZ	ARG	32	-55.820	52.609	-34.305	1.00	20.00
	ATOM	248	NH1	ARG	32	-55.456	52.063	-33.152	1.00	20.00
	ATOM	249	NH2	ARG	32	-55.149	52.327	-35.413	1.00	20.00
55	ATOM	250	N	TYR	33	-61.213	57.250	-30.514	1.00	15.00
	ATOM	251	CA	TYR	33	-61.873	57.632	-29.278	1.00	15.00
	ATOM	252	C	TYR	33	-63.313	58.167	-29.515	1.00	15.00
	ATOM	253	O	TYR	33	-64.243	57.716	-28.859	1.00	15.00
	ATOM	254	CB	TYR	33	-61.016	58.643	-28.585	1.00	15.00
60	ATOM	255	CG	TYR	33	-59.611	58.134	-28.320	1.00	20.00
	ATOM	256	CD1	TYR	33	-58.575	58.331	-29.221	1.00	20.00
	ATOM	257	CD2	TYR	33	-59.328	57.472	-27.131	1.00	20.00
	ATOM	258	CE1	TYR	33	-57.297	57.885	-28.949	1.00	20.00
	ATOM	259	CE2	TYR	33	-58.052	57.024	-26.849	1.00	20.00
65	ATOM	260	CZ	TYR	33	-57.040	57.233	-27.761	1.00	20.00
	ATOM	261	OH	TYR	33	-55.767	56.788	-27.484	1.00	20.00

	ATOM	262	N	CYS	34	-63.488	59.117	-30.449	1.00	15.00
	ATOM	263	CA	CYS	34	-64.801	59.659	-30.766	1.00	15.00
	ATOM	264	C	CYS	34	-65.767	58.497	-31.106	1.00	15.00
	ATOM	265	O	CYS	34	-66.948	58.545	-30.759	1.00	15.00
5	ATOM	266	CB	CYS	34	-64.707	60.685	-31.889	1.00	15.00
	ATOM	267	SG	CYS	34	-63.805	62.207	-31.461	1.00	20.00
	ATOM	268	N	GLN	35	-65.239	57.475	-31.774	1.00	15.00
	ATOM	269	CA	GLN	35	-65.937	56.242	-32.191	1.00	15.00
	ATOM	270	CB	GLN	35	-65.008	55.480	-33.138	1.00	15.00
10	ATOM	271	CG	GLN	35	-65.659	54.433	-34.004	1.00	15.00
	ATOM	272	CD	GLN	35	-66.876	54.985	-34.686	1.00	15.00
	ATOM	273	OE1	GLN	35	-66.803	56.016	-35.351	1.00	15.00
	ATOM	274	NE2	GLN	35	-68.088	54.467	-34.675	1.00	15.00
	ATOM	275	C	GLN	35	-66.278	55.330	-31.016	1.00	15.00
15	ATOM	276	O	GLN	35	-67.404	54.836	-30.905	1.00	15.00
	ATOM	277	N	ALA	36	-65.285	55.146	-30.152	1.00	15.00
	ATOM	278	CA	ALA	36	-65.408	54.359	-28.923	1.00	15.00
	ATOM	279	C	ALA	36	-66.602	54.893	-28.111	1.00	15.00
	ATOM	280	O	ALA	36	-67.459	54.117	-27.681	1.00	15.00
20	ATOM	281	CB	ALA	36	-64.142	54.420	-28.106	1.00	15.00
	ATOM	282	N	SER	37	-66.610	56.232	-27.899	1.00	15.00
	ATOM	283	CA	SER	37	-67.701	56.820	-27.086	1.00	15.00
	ATOM	284	C	SER	37	-69.000	56.736	-27.854	1.00	15.00
	ATOM	285	O	SER	37	-70.064	56.459	-27.275	1.00	15.00
25	ATOM	286	CB	SER	37	-67.357	58.248	-26.632	1.00	15.00
	ATOM	287	OG	SER	37	-66.909	59.048	-27.712	1.00	20.00
	ATOM	288	N	VAL	38	-68.921	56.977	-29.166	1.00	15.00
	ATOM	289	CA	VAL	38	-70.076	56.882	-30.014	1.00	15.00
	ATOM	290	C	VAL	38	-70.770	55.521	-29.876	1.00	15.00
30	ATOM	291	O	VAL	38	-71.989	55.427	-29.641	1.00	15.00
	ATOM	292	CB	VAL	38	-69.675	57.093	-31.504	1.00	15.00
	ATOM	293	CG1	VAL	38	-70.816	56.706	-32.445	1.00	20.00
	ATOM	294	CG2	VAL	38	-69.278	58.546	-31.744	1.00	20.00
	ATOM	295	N	THR	39	-69.966	54.461	-30.061	1.00	15.00
35	ATOM	296	CA	THR	39	-70.482	53.096	-29.987	1.00	15.00
	ATOM	297	C	THR	39	-71.412	52.947	-28.792	1.00	15.00
	ATOM	298	O	THR	39	-72.033	51.895	-28.607	1.00	15.00
	ATOM	299	CB	THR	39	-69.313	52.106	-29.862	1.00	15.00
	ATOM	300	OG1	THR	39	-68.499	52.171	-31.045	1.00	20.00
40	ATOM	301	CG2	THR	39	-69.847	50.706	-29.668	1.00	20.00
	END					0.000	0.000	0.000	0.00	0.00

TABLE 20

111									
5	ATOM	1	C	CYS	1	-69.756	98.907	0.322	1.00101.54
	ATOM	2	O	CYS	1	-69.081	99.101	1.331	1.00102.94
	ATOM	3	CB	CYS	1	-68.777	99.199	-1.962	1.00 96.91
	ATOM	4	SG	CYS	1	-69.633	98.752	-3.506	1.00 92.39
10	ATOM	5	N	CYS	1	-69.207	101.167	-0.527	1.00 93.44
	ATOM	6	CA	CYS	1	-69.685	99.825	-0.884	1.00 98.53
	ATOM	7	N	SER	2	-70.606	97.892	0.139	1.00104.82
	ATOM	8	CA	SER	2	-70.868	96.754	1.009	1.00107.58
15	ATOM	9	CB	SER	2	-71.089	97.133	2.485	1.00109.37
	ATOM	10	OG	SER	2	-70.487	96.196	3.354	1.00110.15
	ATOM	11	C	SER	2	-72.078	95.989	0.410	1.00108.57
	ATOM	12	O	SER	2	-71.941	95.431	-0.681	1.00109.82
20	ATOM	13	N	GLN	3	-73.235	95.942	1.073	1.00108.52
	ATOM	14	CA	GLN	3	-74.362	95.118	0.595	1.00109.56
	ATOM	15	CB	GLN	3	-75.218	95.843	-0.461	1.00108.23
	ATOM	16	CG	GLN	3	-74.627	95.903	-1.853	1.00109.80
25	ATOM	17	CD	GLN	3	-74.110	97.290	-2.104	1.00110.35
	ATOM	18	OE1	GLN	3	-74.838	98.275	-2.037	1.00110.92
	ATOM	19	NE2	GLN	3	-72.861	97.596	-2.416	1.00110.59
	ATOM	20	C	GLN	3	-73.846	93.742	0.105	1.00110.79
30	ATOM	21	O	GLN	3	-74.086	93.328	-1.033	1.00114.50
	ATOM	22	N	ASN	4	-73.129	93.049	1.006	1.00109.87
	ATOM	23	CA	ASN	4	-72.526	91.696	0.886	1.00107.78
	ATOM	24	CB	ASN	4	-73.598	90.633	1.080	1.00109.08
35	ATOM	25	CG	ASN	4	-74.838	90.834	0.214	1.00109.99
	ATOM	26	OD1	ASN	4	-75.620	91.762	0.428	1.00109.73
	ATOM	27	ND2	ASN	4	-75.018	89.946	-0.766	1.00109.43
	ATOM	28	C	ASN	4	-71.718	91.470	-0.396	1.00105.15
40	ATOM	29	O	ASN	4	-72.065	90.635	-1.216	1.00103.42
	ATOM	30	N	GLU	5	-70.624	92.252	-0.576	1.00101.11
	ATOM	31	CA	GLU	5	-69.748	92.135	-1.790	1.00 95.31
	ATOM	32	CB	GLU	5	-70.354	92.907	-2.963	1.00 92.47
45	ATOM	33	CG	GLU	5	-71.871	93.008	-2.964	1.00 86.82
	ATOM	34	CD	GLU	5	-72.288	94.437	-3.248	1.00 85.55
	ATOM	35	OE1	GLU	5	-71.669	95.350	-2.675	1.00 85.72
	ATOM	36	OE2	GLU	5	-73.222	94.621	-4.042	1.00 81.01
50	ATOM	37	C	GLU	5	-68.276	92.590	-1.625	1.00 92.66
	ATOM	38	O	GLU	5	-67.720	92.585	-0.540	1.00 94.43
	ATOM	39	N	TYR	6	-67.689	92.973	-2.759	1.00 85.19
	ATOM	40	CA	TYR	6	-66.295	93.418	-2.833	1.00 78.14
55	ATOM	41	CB	TYR	6	-65.366	92.242	-3.062	1.00 76.59
	ATOM	42	CG	TYR	6	-65.348	91.715	-4.463	1.00 73.94
	ATOM	43	CD1	TYR	6	-64.284	92.059	-5.310	1.00 73.49
	ATOM	44	CE1	TYR	6	-64.225	91.573	-6.601	1.00 73.13
60	ATOM	45	CD2	TYR	6	-66.351	90.880	-4.964	1.00 72.45
	ATOM	46	CE2	TYR	6	-66.297	90.387	-6.276	1.00 72.38
	ATOM	47	CZ	TYR	6	-65.234	90.732	-7.076	1.00 74.15
	ATOM	48	OH	TYR	6	-65.167	90.261	-8.366	1.00 73.32
65	ATOM	49	C	TYR	6	-66.178	94.426	-3.954	1.00 75.43
	ATOM	50	O	TYR	6	-67.043	94.471	-4.849	1.00 76.93
	ATOM	51	N	PHE	7	-65.124	95.237	-3.932	1.00 69.11
	ATOM	52	CA	PHE	7	-64.925	96.212	-4.997	1.00 62.54
70	ATOM	53	CB	PHE	7	-64.530	97.562	-4.389	1.00 61.14
	ATOM	54	CG	PHE	7	-64.206	98.587	-5.441	1.00 61.11
	ATOM	55	CD1	PHE	7	-65.244	99.197	-6.136	1.00 59.90
	ATOM	56	CD2	PHE	7	-62.892	98.921	-5.765	1.00 63.23
75	ATOM	57	CE1	PHE	7	-64.977	100.126	-7.143	1.00 61.45
	ATOM	58	CE2	PHE	7	-62.605	99.850	-6.773	1.00 61.15
	ATOM	59	CZ	PHE	7	-63.647	100.453	-7.463	1.00 61.89
	ATOM	60	C	PHE	7	-63.868	95.730	-6.004	1.00 60.79
80	ATOM	61	O	PHE	7	-62.668	95.911	-5.778	1.00 63.10
	ATOM	62	N	ASP	8	-64.308	95.120	-7.108	1.00 57.06
	ATOM	63	CA	ASP	8	-63.389	94.621	-8.139	1.00 52.20

	ATOM	64	CB	ASP	8	-64.124	93.775	-9.174	1.00	46.52
	ATOM	65	CG	ASP	8	-63.216	93.002	-10.121	1.00	47.00
	ATOM	66	OD1	ASP	8	-62.132	93.534	-10.448	1.00	49.00
	ATOM	67	OD2	ASP	8	-63.596	91.887	-10.531	1.00	48.43
5	ATOM	68	C	ASP	8	-62.695	95.802	-8.830	1.00	53.02
	ATOM	69	O	ASP	8	-63.328	96.562	-9.571	1.00	53.25
	ATOM	70	N	SER	9	-61.395	95.957	-8.578	1.00	55.34
	ATOM	71	CA	SER	9	-60.610	97.041	-9.181	1.00	54.13
	ATOM	72	CB	SER	9	-59.204	97.099	-8.574	1.00	54.65
10	ATOM	73	OG	SER	9	-59.270	97.345	-7.181	1.00	54.09
	ATOM	74	C	SER	9	-60.481	96.887	-10.699	1.00	54.96
	ATOM	75	O	SER	9	-60.309	97.883	-11.421	1.00	54.11
	ATOM	76	N	LEU	10	-60.566	95.657	-11.197	1.00	49.64
	ATOM	77	CA	LEU	10	-60.477	95.470	-12.630	1.00	46.11
15	ATOM	78	CB	LEU	10	-60.454	93.995	-13.003	1.00	40.39
	ATOM	79	CG	LEU	10	-60.157	93.681	-14.453	1.00	34.58
	ATOM	80	CD1	LEU	10	-58.716	94.067	-14.797	1.00	33.04
	ATOM	81	CD2	LEU	10	-60.417	92.204	-14.758	1.00	33.82
	ATOM	82	C	LEU	10	-61.666	96.194	-13.262	1.00	49.82
20	ATOM	83	O	LEU	10	-61.513	96.913	-14.235	1.00	50.94
	ATOM	84	N	LEU	11	-62.858	95.992	-12.700	1.00	51.20
	ATOM	85	CA	LEU	11	-64.064	96.618	-13.238	1.00	51.39
	ATOM	86	CB	LEU	11	-65.289	95.744	-12.949	1.00	47.82
	ATOM	87	CG	LEU	11	-65.267	94.286	-13.431	1.00	46.28
25	ATOM	88	CD1	LEU	11	-66.589	93.602	-13.102	1.00	49.44
	ATOM	89	CD2	LEU	11	-64.988	94.204	-14.925	1.00	44.53
	ATOM	90	C	LEU	11	-64.332	98.040	-12.732	1.00	53.87
	ATOM	91	O	LEU	11	-65.034	98.808	-13.377	1.00	50.27
	ATOM	92	N	HIS	12	-63.780	98.386	-11.563	1.00	58.57
30	ATOM	93	CA	HIS	12	-63.990	99.682	-10.951	1.00	64.33
	ATOM	94	CB	HIS	12	-63.637	100.824	-11.902	1.00	65.68
	ATOM	95	CG	HIS	12	-62.125	100.890	-12.238	1.00	69.20
	ATOM	96	CD2	HIS	12	-61.036	100.544	-11.509	1.00	69.34
	ATOM	97	ND1	HIS	12	-61.646	101.325	-13.456	1.00	71.64
35	ATOM	98	CE1	HIS	12	-60.324	101.235	-13.467	1.00	72.29
	ATOM	99	NE2	HIS	12	-59.930	100.764	-12.297	1.00	70.37
	ATOM	100	C	HIS	12	-65.453	99.852	-10.568	1.00	66.69
	ATOM	101	O	HIS	12	-66.063	100.909	-10.759	1.00	67.11
	ATOM	102	N	ALA	13	-66.015	98.770	-10.036	1.00	69.57
40	ATOM	103	CA	ALA	13	-67.416	98.727	-9.640	1.00	71.62
	ATOM	104	CB	ALA	13	-68.302	98.359	-10.807	1.00	69.66
	ATOM	105	C	ALA	13	-67.570	97.718	-8.506	1.00	73.99
	ATOM	106	O	ALA	13	-66.577	97.145	-8.041	1.00	75.45
	ATOM	107	N	CYS	14	-68.806	97.482	-8.070	1.00	75.75
45	ATOM	108	CA	CYS	14	-69.036	96.552	-6.971	1.00	77.01
	ATOM	109	C	CYS	14	-69.732	95.266	-7.367	1.00	75.86
	ATOM	110	O	CYS	14	-70.863	95.258	-7.863	1.00	73.08
	ATOM	111	CB	CYS	14	-69.787	97.267	-5.847	1.00	80.96
	ATOM	112	SG	CYS	14	-68.726	97.877	-4.501	1.00	86.58
50	ATOM	113	N	ILE	15	-69.002	94.159	-7.115	1.00	75.89
	ATOM	114	CA	ILE	15	-69.494	92.836	-7.466	1.00	77.02
	ATOM	115	CB	ILE	15	-68.459	92.064	-8.318	1.00	76.68
	ATOM	116	CG2	ILE	15	-68.983	90.673	-8.660	1.00	76.37
	ATOM	117	CG1	ILE	15	-68.109	92.833	-9.604	1.00	76.28
55	ATOM	118	CD1	ILE	15	-69.257	92.981	-10.580	1.00	74.77
	ATOM	119	C	ILE	15	-69.892	91.980	-6.241	1.00	78.19
	ATOM	120	O	ILE	15	-69.196	91.897	-5.240	1.00	77.54
	ATOM	121	N	PRO	16	-71.073	91.329	-6.422	1.00	80.09
	ATOM	122	CD	PRO	16	-72.234	92.040	-7.004	1.00	79.64
60	ATOM	123	CA	PRO	16	-71.578	90.386	-5.379	1.00	82.16
	ATOM	124	CB	PRO	16	-72.890	89.912	-5.980	1.00	80.00
	ATOM	125	CG	PRO	16	-73.388	91.147	-6.648	1.00	80.17
	ATOM	126	C	PRO	16	-70.568	89.371	-4.930	1.00	84.28
	ATOM	127	O	PRO	16	-70.028	88.654	-5.771	1.00	84.12
65	ATOM	128	N	CYS	17	-70.274	89.279	-3.629	1.00	87.37
	ATOM	129	CA	CYS	17	-69.298	88.307	-3.162	1.00	88.33

	ATOM	130	CB	CYS	17	-69.296	88.245	-1.637	1.00	86.90
	ATOM	131	SG	CYS	17	-68.385	89.595	-0.828	1.00	84.56
	ATOM	132	C	CYS	17	-69.567	86.913	-3.741	1.00	89.64
	ATOM	133	O	CYS	17	-68.631	86.184	-4.107	1.00	89.77
5	ATOM	134	N	GLN	18	-70.850	86.556	-3.812	1.00	91.08
	ATOM	135	CA	GLN	18	-71.259	85.297	-4.363	1.00	91.16
	ATOM	136	CB	GLN	18	-72.721	85.312	-4.810	1.00	92.68
	ATOM	137	CG	GLN	18	-73.633	86.048	-3.834	1.00	98.31
	ATOM	138	CD	GLN	18	-74.942	86.558	-4.448	1.00	101.37
10	ATOM	139	OE1	GLN	18	-74.941	87.480	-5.255	1.00	102.74
	ATOM	140	NE2	GLN	18	-76.150	86.088	-4.199	1.00	103.76
	ATOM	141	C	GLN	18	-70.362	84.974	-5.561	1.00	89.18
	ATOM	142	O	GLN	18	-69.718	83.924	-5.621	1.00	89.64
	ATOM	143	N	LEU	19	-70.358	85.914	-6.508	1.00	87.56
15	ATOM	144	CA	LEU	19	-69.620	85.805	-7.754	1.00	85.33
	ATOM	145	CB	LEU	19	-69.657	87.116	-8.486	1.00	79.98
	ATOM	146	CG	LEU	19	-70.635	87.136	-9.641	1.00	77.91
	ATOM	147	CD1	LEU	19	-70.906	88.562	-10.091	1.00	78.13
	ATOM	148	CD2	LEU	19	-70.124	86.296	-10.799	1.00	72.50
20	ATOM	149	C	LEU	19	-68.179	85.344	-7.612	1.00	86.05
	ATOM	150	O	LEU	19	-67.728	84.476	-8.375	1.00	88.06
	ATOM	151	N	ARG	20	-67.457	85.888	-6.662	1.00	85.11
	ATOM	152	CA	ARG	20	-66.046	85.510	-6.517	1.00	83.33
	ATOM	153	CB	ARG	20	-65.273	86.625	-5.817	1.00	78.93
25	ATOM	154	CG	ARG	20	-64.408	87.475	-6.742	1.00	73.68
	ATOM	155	CD	ARG	20	-63.253	86.682	-7.327	1.00	67.23
	ATOM	156	NE	ARG	20	-61.958	87.183	-6.865	1.00	63.84
	ATOM	157	CZ	ARG	20	-61.596	88.467	-6.896	1.00	64.39
	ATOM	158	NH1	ARG	20	-62.445	89.374	-7.371	1.00	66.99
30	ATOM	159	NH2	ARG	20	-60.399	88.829	-6.462	1.00	60.63
	ATOM	160	C	ARG	20	-65.841	84.178	-5.793	1.00	84.21
	ATOM	161	O	ARG	20	-64.740	83.917	-5.310	1.00	86.28
	ATOM	162	N	CYS	21	-66.871	83.347	-5.715	1.00	85.02
	ATOM	163	CA	CYS	21	-66.685	82.057	-5.060	1.00	83.51
35	ATOM	164	CB	CYS	21	-67.752	81.877	-3.971	1.00	81.05
	ATOM	165	SG	CYS	21	-67.750	83.175	-2.693	1.00	73.08
	ATOM	166	C	CYS	21	-66.668	80.870	-6.057	1.00	84.26
	ATOM	167	O	CYS	21	-67.597	80.076	-6.052	1.00	84.48
	ATOM	168	N	SER	22	-65.615	80.748	-6.942	1.00	84.61
40	ATOM	169	CA	SER	22	-65.437	79.663	-7.955	1.00	85.07
	ATOM	170	C	SER	22	-64.254	78.782	-7.584	1.00	85.65
	ATOM	171	O	SER	22	-63.639	78.122	-8.419	1.00	86.77
	ATOM	172	CB	SER	22	-65.187	80.232	-9.359	1.00	20.00
	ATOM	173	OG	SER	22	-63.810	80.484	-9.562	1.00	20.00
45	ATOM	174	N	SER	23	-63.993	78.811	-6.306	1.00	84.19
	ATOM	175	CA	SER	23	-62.949	77.998	-5.725	1.00	82.44
	ATOM	176	C	SER	23	-61.656	78.769	-5.435	1.00	84.32
	ATOM	177	O	SER	23	-61.585	79.437	-4.402	1.00	81.24
	ATOM	178	CB	SER	23	-62.721	76.789	-6.633	1.00	78.00
50	ATOM	179	OG	SER	23	-61.355	76.424	-6.676	1.00	20.00
	ATOM	180	N	ASN	24	-60.661	78.709	-6.273	1.00	87.06
	ATOM	181	CA	ASN	24	-59.496	79.484	-5.896	1.00	88.17
	ATOM	182	C	ASN	24	-59.886	80.968	-5.833	1.00	88.45
	ATOM	183	O	ASN	24	-59.028	81.808	-5.625	1.00	84.89
55	ATOM	184	CB	ASN	24	-58.384	79.372	-6.916	0.00	86.84
	ATOM	185	CG	ASN	24	-58.017	77.963	-7.210	0.00	85.55
	ATOM	186	OD1	ASN	24	-58.189	77.089	-6.364	1.00	20.00
	ATOM	187	ND2	ASN	24	-57.501	77.709	-8.406	1.00	20.00
	ATOM	188	N	THR	25	-61.195	81.281	-6.023	1.00	89.49
60	ATOM	189	CA	THR	25	-61.570	82.706	-6.136	1.00	91.86
	ATOM	190	C	THR	25	-62.260	83.454	-4.944	1.00	95.75
	ATOM	191	O	THR	25	-62.999	84.433	-5.141	1.00	96.59
	ATOM	192	CB	THR	25	-62.349	82.868	-7.423	0.00	91.23
	ATOM	193	OG1	THR	25	-61.567	82.370	-8.509	1.00	20.00
65	ATOM	194	CG2	THR	25	-62.689	84.326	-7.643	1.00	20.00
	ATOM	195	N	PRO	26	-62.013	82.970	-3.705	1.00	98.41

	ATOM	196	CA	PRO	26	-62.431	83.691	-2.382	1.00	96.39
	ATOM	197	C	PRO	26	-61.676	85.062	-1.819	1.00	96.88
	ATOM	198	O	PRO	26	-60.679	84.885	-1.114	1.00	94.52
	ATOM	199	CB	PRO	26	-62.321	82.539	-1.402	0.00	95.29
5	ATOM	200	CG	PRO	26	-62.797	81.345	-2.175	0.00	92.34
	ATOM	201	CD	PRO	26	-62.594	81.630	-3.629	0.00	89.50
	ATOM	202	N	PRO	27	-62.090	86.393	-2.122	1.00	97.72
	ATOM	203	CA	PRO	27	-61.471	87.711	-1.527	1.00	98.94
	ATOM	204	C	PRO	27	-61.478	88.055	0.024	1.00	100.71
10	ATOM	205	O	PRO	27	-62.222	87.407	0.757	1.00	100.56
	ATOM	206	CB	PRO	27	-62.106	88.769	-2.402	0.00	97.33
	ATOM	207	CG	PRO	27	-62.134	88.106	-3.740	0.00	94.51
	ATOM	208	CD	PRO	27	-62.165	86.632	-3.552	0.00	92.58
	ATOM	209	N	LEU	28	-60.680	89.050	0.581	1.00	15.00
15	ATOM	210	CA	LEU	28	-60.934	89.368	2.025	1.00	15.00
	ATOM	211	C	LEU	28	-62.443	89.590	1.970	1.00	15.00
	ATOM	212	O	LEU	28	-63.096	89.131	1.032	1.00	15.00
	ATOM	213	CB	LEU	28	-60.176	90.635	2.550	1.00	15.00
	ATOM	214	CG	LEU	28	-59.159	90.459	3.721	1.00	15.00
20	ATOM	215	CD1	LEU	28	-59.667	91.157	4.961	1.00	15.00
	ATOM	216	CD2	LEU	28	-58.900	88.978	3.994	1.00	15.00
	ATOM	217	N	THR	29	-62.971	90.312	2.944	1.00	15.00
	ATOM	218	CA	THR	29	-64.391	90.642	2.905	1.00	15.00
	ATOM	219	C	THR	29	-65.147	89.651	2.026	1.00	15.00
25	ATOM	220	O	THR	29	-66.213	89.955	1.516	1.00	15.00
	ATOM	221	CB	THR	29	-64.578	92.081	2.409	1.00	15.00
	ATOM	222	OG1	THR	29	-64.799	92.933	3.529	1.00	20.00
	ATOM	223	CG2	THR	29	-65.763	92.149	1.454	1.00	20.00
	ATOM	224	N	CYS	30	-64.551	88.455	1.887	1.00	15.00
30	ATOM	225	CA	CYS	30	-65.164	87.454	1.035	1.00	15.00
	ATOM	226	C	CYS	30	-65.052	86.072	1.665	1.00	15.00
	ATOM	227	O	CYS	30	-65.908	85.231	1.445	1.00	15.00
	ATOM	228	CB	CYS	30	-64.583	87.524	-0.370	1.00	15.00
	ATOM	229	SG	CYS	30	-65.125	88.957	-1.353	1.00	20.00
35	ATOM	230	N	GLN	31	-63.997	85.859	2.452	1.00	15.00
	ATOM	231	CA	GLN	31	-63.848	84.588	3.120	1.00	15.00
	ATOM	232	C	GLN	31	-65.049	84.288	4.021	1.00	15.00
	ATOM	233	O	GLN	31	-65.643	83.206	3.969	1.00	15.00
	ATOM	234	CB	GLN	31	-62.568	84.545	3.915	1.00	15.00
40	ATOM	235	CG	GLN	31	-61.396	84.109	3.046	1.00	20.00
	ATOM	236	CD	GLN	31	-60.110	83.952	3.832	1.00	20.00
	ATOM	237	OE1	GLN	31	-60.130	83.825	5.056	1.00	20.00
	ATOM	238	NE2	GLN	31	-58.878	83.932	3.333	1.00	20.00
	ATOM	239	N	ARG	32	-65.403	85.245	4.832	1.00	15.00
45	ATOM	240	CA	ARG	32	-66.545	85.143	5.716	1.00	15.00
	ATOM	241	C	ARG	32	-67.834	84.865	4.948	1.00	15.00
	ATOM	242	O	ARG	32	-68.680	84.064	5.381	1.00	15.00
	ATOM	243	CB	ARG	32	-66.642	86.412	6.589	1.00	15.00
	ATOM	244	CG	ARG	32	-65.438	86.655	7.504	1.00	15.00
50	ATOM	245	CD	ARG	32	-65.671	87.884	8.377	1.00	15.00
	ATOM	246	NE	ARG	32	-64.524	88.207	9.217	1.00	20.00
	ATOM	247	CZ	ARG	32	-64.507	89.194	10.105	1.00	20.00
	ATOM	248	NH1	ARG	32	-65.581	89.956	10.268	1.00	20.00
	ATOM	249	NH2	ARG	32	-63.420	89.417	10.831	1.00	20.00
55	ATOM	250	N	TYR	33	-67.945	85.530	3.801	1.00	15.00
	ATOM	251	CA	TYR	33	-69.103	85.364	2.940	1.00	15.00
	ATOM	252	C	TYR	33	-69.194	83.933	2.337	1.00	15.00
	ATOM	253	O	TYR	33	-70.243	83.308	2.415	1.00	15.00
	ATOM	254	CB	TYR	33	-69.045	86.410	1.872	1.00	15.00
60	ATOM	255	CG	TYR	33	-68.981	87.817	2.437	1.00	20.00
	ATOM	256	CD1	TYR	33	-67.776	88.453	2.693	1.00	20.00
	ATOM	257	CD2	TYR	33	-70.156	88.516	2.686	1.00	20.00
	ATOM	258	CE1	TYR	33	-67.738	89.744	3.180	1.00	20.00
	ATOM	259	CE2	TYR	33	-70.129	89.809	3.171	1.00	20.00
65	ATOM	260	CZ	TYR	33	-68.918	90.419	3.416	1.00	20.00
	ATOM	261	OH	TYR	33	-68.886	91.707	3.900	1.00	20.00

	ATOM	262	N	CYS	34	-68.098	83.426	1.747	1.00	15.00
	ATOM	263	CA	CYS	34	-68.073	82.086	1.181	1.00	15.00
	ATOM	264	C	CYS	34	-68.530	81.067	2.255	1.00	15.00
	ATOM	265	O	CYS	34	-69.216	80.094	1.941	1.00	15.00
5	ATOM	266	CB	CYS	34	-66.694	81.759	0.619	1.00	15.00
	ATOM	267	SG	CYS	34	-66.206	82.749	-0.828	1.00	20.00
	ATOM	268	N	GLN	35	-68.142	81.321	3.502	1.00	15.00
	ATOM	269	CA	GLN	35	-68.467	80.524	4.704	1.00	15.00
	ATOM	270	CB	GLN	35	-67.606	81.049	5.855	1.00	15.00
10	ATOM	271	CG	GLN	35	-67.457	80.131	7.041	1.00	15.00
	ATOM	272	CD	GLN	35	-67.080	78.747	6.602	1.00	15.00
	ATOM	273	OE1	GLN	35	-66.103	78.569	5.878	1.00	15.00
	ATOM	274	NE2	GLN	35	-67.690	77.624	6.925	1.00	15.00
	ATOM	275	C	GLN	35	-69.933	80.639	5.110	1.00	15.00
15	ATOM	276	O	GLN	35	-70.592	79.632	5.388	1.00	15.00
	ATOM	277	N	ALA	36	-70.411	81.878	5.115	1.00	15.00
	ATOM	278	CA	ALA	36	-71.804	82.214	5.415	1.00	15.00
	ATOM	279	C	ALA	36	-72.721	81.396	4.488	1.00	15.00
	ATOM	280	O	ALA	36	-73.667	80.758	4.955	1.00	15.00
20	ATOM	281	CB	ALA	36	-72.059	83.691	5.243	1.00	15.00
	ATOM	282	N	SER	37	-72.421	81.459	3.167	1.00	15.00
	ATOM	283	CA	SER	37	-73.284	80.736	2.203	1.00	15.00
	ATOM	284	C	SER	37	-73.091	79.248	2.376	1.00	15.00
	ATOM	285	O	SER	37	-74.056	78.469	2.300	1.00	15.00
25	ATOM	286	CB	SER	37	-73.041	81.215	0.763	1.00	15.00
	ATOM	287	OG	SER	37	-71.662	81.236	0.442	1.00	20.00
	ATOM	288	N	VAL	38	-71.839	78.842	2.607	1.00	15.00
	ATOM	289	CA	VAL	38	-71.532	77.458	2.836	1.00	15.00
	ATOM	290	C	VAL	38	-72.382	76.868	3.968	1.00	15.00
30	ATOM	291	O	VAL	38	-73.035	75.819	3.820	1.00	15.00
	ATOM	292	CB	VAL	38	-70.026	77.288	3.198	1.00	15.00
	ATOM	293	CG1	VAL	38	-69.739	75.885	3.732	1.00	20.00
	ATOM	294	CG2	VAL	38	-69.154	77.565	1.977	1.00	20.00
	ATOM	295	N	THR	39	-72.333	77.555	5.122	1.00	15.00
35	ATOM	296	CA	THR	39	-73.069	77.107	6.302	1.00	15.00
	ATOM	297	C	THR	39	-74.476	76.676	5.915	1.00	15.00
	ATOM	298	O	THR	39	-75.230	76.171	6.752	1.00	15.00
	ATOM	299	CB	THR	39	-73.139	78.246	7.332	1.00	15.00
	ATOM	300	OG1	THR	39	-71.812	78.573	7.777	1.00	20.00
40	ATOM	301	CG2	THR	39	-73.997	77.825	8.502	1.00	20.00
	END					-94.358	153.908	54.620	0.00	0.00

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	111								
	ATOM	1	C	CYS	1	-82.437	88.466	-46.488	1.00101.54
5	ATOM	2	O	CYS	1	-83.020	89.459	-46.916	1.00102.94
	ATOM	3	CB	CYS	1	-80.014	88.556	-45.871	1.00 96.91
	ATOM	4	SG	CYS	1	-79.115	87.201	-45.053	1.00 92.39
	ATOM	5	N	CYS	1	-80.705	88.677	-48.243	1.00 93.44
	ATOM	6	CA	CYS	1	-81.034	88.100	-46.934	1.00 98.53
10	ATOM	7	N	SER	2	-82.926	87.603	-45.591	1.00104.82
	ATOM	8	CA	SER	2	-84.179	87.669	-44.851	1.00107.58
	ATOM	9	CB	SER	2	-85.404	87.996	-45.725	1.00109.37
	ATOM	10	OG	SER	2	-86.294	88.868	-45.061	1.00110.15
	ATOM	11	C	SER	2	-84.337	86.323	-44.097	1.00108.57
15	ATOM	12	O	SER	2	-83.540	86.056	-43.195	1.00109.82
	ATOM	13	N	GLN	3	-85.320	85.483	-44.426	1.00108.52
	ATOM	14	CA	GLN	3	-85.576	84.256	-43.645	1.00109.56
	ATOM	15	CB	GLN	3	-84.684	83.079	-44.083	1.00108.23
	ATOM	16	CG	GLN	3	-83.251	83.129	-43.598	1.00109.80
20	ATOM	17	CD	GLN	3	-82.362	83.525	-44.742	1.00110.35
	ATOM	18	OE1	GLN	3	-82.312	82.873	-45.779	1.00110.92
	ATOM	19	NE2	GLN	3	-81.562	84.579	-44.760	1.00110.59
	ATOM	20	C	GLN	3	-85.468	84.558	-42.130	1.00110.79
	ATOM	21	O	GLN	3	-84.705	83.923	-41.395	1.00114.50
25	ATOM	22	N	ASN	4	-86.261	85.548	-41.688	1.00109.87
	ATOM	23	CA	ASN	4	-86.438	86.064	-40.305	1.00107.78
	ATOM	24	CB	ASN	4	-87.347	85.131	-39.518	1.00109.08
	ATOM	25	CG	ASN	4	-86.936	83.663	-39.581	1.00109.99
	ATOM	26	OD1	ASN	4	-87.053	83.013	-40.620	1.00109.73
30	ATOM	27	ND2	ASN	4	-86.460	83.140	-38.448	1.00109.43
	ATOM	28	C	ASN	4	-85.131	86.356	-39.561	1.00105.15
	ATOM	29	O	ASN	4	-84.831	85.734	-38.554	1.00103.42
	ATOM	30	N	GLU	5	-84.327	87.313	-40.088	1.00101.11
	ATOM	31	CA	GLU	5	-83.017	87.693	-39.460	1.00 95.31
35	ATOM	32	CB	GLU	5	-81.922	86.707	-39.870	1.00 92.47
	ATOM	33	CG	GLU	5	-82.393	85.292	-40.163	1.00 86.82
	ATOM	34	CD	GLU	5	-81.774	84.805	-41.458	1.00 85.55
	ATOM	35	OE1	GLU	5	-81.740	85.591	-42.420	1.00 85.72
	ATOM	36	OE2	GLU	5	-81.330	83.647	-41.488	1.00 81.01
40	ATOM	37	C	GLU	5	-82.504	89.127	-39.747	1.00 92.66
	ATOM	38	O	GLU	5	-83.265	90.036	-40.029	1.00 94.43
	ATOM	39	N	TYR	6	-81.182	89.267	-39.650	1.00 85.19
	ATOM	40	CA	TYR	6	-80.491	90.542	-39.858	1.00 78.14
	ATOM	41	CB	TYR	6	-80.400	91.323	-38.561	1.00 76.59
45	ATOM	42	CG	TYR	6	-79.362	90.833	-37.600	1.00 73.94
	ATOM	43	CD1	TYR	6	-78.143	91.521	-37.500	1.00 73.49
	ATOM	44	CE1	TYR	6	-77.173	91.110	-36.609	1.00 73.13
	ATOM	45	CD2	TYR	6	-79.560	89.715	-36.786	1.00 72.45
	ATOM	46	CE2	TYR	6	-78.575	89.291	-35.881	1.00 72.38
50	ATOM	47	CZ	TYR	6	-77.398	89.995	-35.798	1.00 74.15
	ATOM	48	OH	TYR	6	-76.420	89.591	-34.919	1.00 73.32
	ATOM	49	C	TYR	6	-79.114	90.250	-40.410	1.00 75.43
	ATOM	50	O	TYR	6	-78.608	89.121	-40.268	1.00 76.93
	ATOM	51	N	PHE	7	-78.489	91.243	-41.036	1.00 69.11
55	ATOM	52	CA	PHE	7	-77.145	91.048	-41.566	1.00 62.54
	ATOM	53	CB	PHE	7	-77.060	91.639	-42.977	1.00 61.14
	ATOM	54	CG	PHE	7	-75.667	91.566	-43.542	1.00 61.11
	ATOM	55	CD1	PHE	7	-75.192	90.348	-44.017	1.00 59.90
	ATOM	56	CD2	PHE	7	-74.825	92.675	-43.573	1.00 63.23
60	ATOM	57	CE1	PHE	7	-73.893	90.237	-44.514	1.00 61.45
	ATOM	58	CE2	PHE	7	-73.518	92.583	-44.068	1.00 61.15
	ATOM	59	CZ	PHE	7	-73.050	91.364	-44.538	1.00 61.89
	ATOM	60	C	PHE	7	-76.086	91.670	-40.642	1.00 60.79
	ATOM	61	O	PHE	7	-75.815	92.871	-40.729	1.00 63.10
65	ATOM	62	N	ASP	8	-75.489	90.860	-39.763	1.00 57.06
	ATOM	63	CA	ASP	8	-74.462	91.344	-38.833	1.00 52.20

	ATOM	64	CB	ASP	8	-74.108	90.283	-37.795	1.00	46.52
	ATOM	65	CG	ASP	8	-73.255	90.787	-36.639	1.00	47.00
	ATOM	66	OD1	ASP	8	-72.416	91.682	-36.884	1.00	49.00
	ATOM	67	OD2	ASP	8	-73.424	90.282	-35.511	1.00	48.43
5	ATOM	68	C	ASP	8	-73.204	91.746	-39.615	1.00	53.02
	ATOM	69	O	ASP	8	-72.498	90.891	-40.161	1.00	53.25
	ATOM	70	N	SER	9	-72.932	93.050	-39.672	1.00	55.34
	ATOM	71	CA	SER	9	-71.755	93.568	-40.381	1.00	54.13
	ATOM	72	CB	SER	9	-71.792	95.098	-40.453	1.00	54.65
10	ATOM	73	OG	SER	9	-72.940	95.539	-41.155	1.00	54.09
	ATOM	74	C	SER	9	-70.444	93.142	-39.715	1.00	54.96
	ATOM	75	O	SER	9	-69.401	93.045	-40.383	1.00	54.11
	ATOM	76	N	LEU	10	-70.479	92.879	-38.412	1.00	49.64
	ATOM	77	CA	LEU	10	-69.267	92.446	-37.749	1.00	46.11
15	ATOM	78	CB	LEU	10	-69.462	92.328	-36.244	1.00	40.39
	ATOM	79	CG	LEU	10	-68.211	92.082	-35.430	1.00	34.58
	ATOM	80	CD1	LEU	10	-67.290	93.303	-35.486	1.00	33.04
	ATOM	81	CD2	LEU	10	-68.561	91.725	-33.983	1.00	33.82
	ATOM	82	C	LEU	10	-68.856	91.110	-38.370	1.00	49.82
20	ATOM	83	O	LEU	10	-67.699	90.904	-38.698	1.00	50.94
	ATOM	84	N	LEU	11	-69.818	90.201	-38.525	1.00	51.20
	ATOM	85	CA	LEU	11	-69.530	88.884	-39.090	1.00	51.39
	ATOM	86	CB	LEU	11	-70.505	87.842	-38.530	1.00	47.82
	ATOM	87	CG	LEU	11	-70.599	87.685	-37.006	1.00	46.28
25	ATOM	88	CD1	LEU	11	-71.574	86.568	-36.650	1.00	49.44
	ATOM	89	CD2	LEU	11	-69.233	87.406	-36.395	1.00	44.53
	ATOM	90	C	LEU	11	-69.552	88.820	-40.621	1.00	53.87
	ATOM	91	O	LEU	11	-68.950	87.935	-41.216	1.00	50.27
	ATOM	92	N	HIS	12	-70.261	89.756	-41.260	1.00	58.57
30	ATOM	93	CA	HIS	12	-70.402	89.786	-42.702	1.00	64.33
	ATOM	94	CB	HIS	12	-69.046	89.775	-43.405	1.00	65.68
	ATOM	95	CG	HIS	12	-68.222	91.065	-43.157	1.00	69.20
	ATOM	96	CD2	HIS	12	-68.616	92.342	-42.933	1.00	69.34
	ATOM	97	ND1	HIS	12	-66.845	91.074	-43.095	1.00	71.64
35	ATOM	98	CE1	HIS	12	-66.424	92.302	-42.834	1.00	72.29
	ATOM	99	NE2	HIS	12	-67.480	93.091	-42.731	1.00	70.37
	ATOM	100	C	HIS	12	-71.165	88.559	-43.180	1.00	66.69
	ATOM	101	O	HIS	12	-70.825	87.923	-44.184	1.00	67.11
	ATOM	102	N	ALA	13	-72.204	88.223	-42.421	1.00	69.57
40	ATOM	103	CA	ALA	13	-73.035	87.059	-42.697	1.00	71.62
	ATOM	104	CB	ALA	13	-72.447	85.811	-42.081	1.00	69.66
	ATOM	105	C	ALA	13	-74.436	87.321	-42.152	1.00	73.99
	ATOM	106	O	ALA	13	-74.713	88.413	-41.641	1.00	75.45
	ATOM	107	N	CYS	14	-75.315	86.324	-42.239	1.00	75.75
45	ATOM	108	CA	CYS	14	-76.683	86.502	-41.766	1.00	77.01
	ATOM	109	C	CYS	14	-77.031	85.707	-40.526	1.00	75.86
	ATOM	110	O	CYS	14	-76.981	84.474	-40.501	1.00	73.08
	ATOM	111	CB	CYS	14	-77.659	86.210	-42.908	1.00	80.96
	ATOM	112	SG	CYS	14	-78.257	87.686	-43.785	1.00	86.58
50	ATOM	113	N	ILE	15	-77.402	86.476	-39.479	1.00	75.89
	ATOM	114	CA	ILE	15	-77.735	85.887	-38.192	1.00	77.02
	ATOM	115	CB	ILE	15	-76.921	86.543	-37.051	1.00	76.68
	ATOM	116	CG2	ILE	15	-77.296	85.928	-35.707	1.00	76.37
	ATOM	117	CG1	ILE	15	-75.408	86.408	-37.296	1.00	76.28
55	ATOM	118	CD1	ILE	15	-74.889	84.986	-37.260	1.00	74.77
	ATOM	119	C	ILE	15	-79.242	85.954	-37.853	1.00	78.19
	ATOM	120	O	ILE	15	-79.910	86.965	-38.017	1.00	77.54
	ATOM	121	N	PRO	16	-79.713	84.786	-37.339	1.00	80.09
	ATOM	122	CD	PRO	16	-79.341	83.495	-37.962	1.00	79.64
60	ATOM	123	CA	PRO	16	-81.129	84.687	-36.872	1.00	82.16
	ATOM	124	CB	PRO	16	-81.215	83.246	-36.402	1.00	80.00
	ATOM	125	CG	PRO	16	-80.358	82.544	-37.399	1.00	80.17
	ATOM	126	C	PRO	16	-81.544	85.789	-35.941	1.00	84.28
	ATOM	127	O	PRO	16	-80.887	85.988	-34.921	1.00	84.12
65	ATOM	128	N	CYS	17	-82.612	86.532	-36.250	1.00	87.37
	ATOM	129	CA	CYS	17	-83.040	87.608	-35.370	1.00	88.33

	ATOM	130	CB	CYS	17	-84.390	88.158	-35.819	1.00	86.90
	ATOM	131	SG	CYS	17	-84.307	89.304	-37.229	1.00	84.56
	ATOM	132	C	CYS	17	-83.124	87.144	-33.911	1.00	89.64
	ATOM	133	O	CYS	17	-82.752	87.883	-32.985	1.00	89.77
5	ATOM	134	N	GLN	18	-83.621	85.921	-33.722	1.00	91.08
	ATOM	135	CA	GLN	18	-83.728	85.338	-32.417	1.00	91.16
	ATOM	136	CB	GLN	18	-83.824	83.812	-32.474	1.00	92.68
	ATOM	137	CG	GLN	18	-84.716	83.316	-33.605	1.00	98.31
	ATOM	138	CD	GLN	18	-84.438	81.874	-34.049	1.00	101.37
10	ATOM	139	OE1	GLN	18	-83.405	81.587	-34.642	1.00	102.74
	ATOM	140	NE2	GLN	18	-85.229	80.836	-33.853	1.00	103.76
	ATOM	141	C	GLN	18	-82.498	85.741	-31.598	1.00	89.18
	ATOM	142	O	GLN	18	-82.606	86.318	-30.514	1.00	89.64
	ATOM	143	N	LEU	19	-81.336	85.407	-32.161	1.00	87.56
15	ATOM	144	CA	LEU	19	-80.041	85.646	-31.547	1.00	85.33
	ATOM	145	CB	LEU	19	-78.947	85.352	-32.532	1.00	79.98
	ATOM	146	CG	LEU	19	-78.261	84.023	-32.295	1.00	77.91
	ATOM	147	CD1	LEU	19	-77.450	83.612	-33.513	1.00	78.13
	ATOM	148	CD2	LEU	19	-77.381	84.080	-31.059	1.00	72.50
20	ATOM	149	C	LEU	19	-79.847	87.040	-30.975	1.00	86.05
	ATOM	150	O	LEU	19	-79.341	87.184	-29.851	1.00	88.06
	ATOM	151	N	ARG	20	-80.238	88.057	-31.704	1.00	85.11
	ATOM	152	CA	ARG	20	-80.026	89.424	-31.214	1.00	83.33
	ATOM	153	CB	ARG	20	-79.978	90.401	-32.386	1.00	78.93
25	ATOM	154	CG	ARG	20	-78.577	90.877	-32.759	1.00	73.68
	ATOM	155	CD	ARG	20	-77.964	91.741	-31.673	1.00	67.23
	ATOM	156	NE	ARG	20	-77.753	93.118	-32.123	1.00	63.84
	ATOM	157	CZ	ARG	20	-77.145	93.448	-33.265	1.00	64.39
	ATOM	158	NH1	ARG	20	-76.691	92.487	-34.065	1.00	66.99
30	ATOM	159	NH2	ARG	20	-76.992	94.721	-33.590	1.00	60.63
	ATOM	160	C	ARG	20	-81.064	89.873	-30.184	1.00	84.21
	ATOM	161	O	ARG	20	-81.210	91.073	-29.956	1.00	86.28
	ATOM	162	N	CYS	21	-81.775	88.938	-29.569	1.00	85.02
	ATOM	163	CA	CYS	21	-82.745	89.344	-28.558	1.00	83.51
35	ATOM	164	CB	CYS	21	-84.117	88.741	-28.893	1.00	81.05
	ATOM	165	SG	CYS	21	-84.765	89.206	-30.530	1.00	73.08
	ATOM	166	C	CYS	21	-82.295	88.997	-27.116	1.00	84.26
	ATOM	167	O	CYS	21	-82.896	88.130	-26.498	1.00	84.48
	ATOM	168	N	SER	22	-81.215	89.660	-26.570	1.00	84.61
40	ATOM	169	CA	SER	22	-80.660	89.458	-25.195	1.00	85.07
	ATOM	170	C	SER	22	-80.902	90.693	-24.340	1.00	85.65
	ATOM	171	O	SER	22	-80.204	90.964	-23.366	1.00	86.77
	ATOM	172	CB	SER	22	-79.149	89.188	-25.226	1.00	20.00
	ATOM	173	OG	SER	22	-78.420	90.400	-25.213	1.00	20.00
45	ATOM	174	N	SER	23	-81.918	91.397	-24.759	1.00	84.19
	ATOM	175	CA	SER	23	-82.364	92.579	-24.056	1.00	82.44
	ATOM	176	C	SER	23	-81.908	93.891	-24.703	1.00	84.32
	ATOM	177	O	SER	23	-82.545	94.331	-25.661	1.00	81.24
	ATOM	178	CB	SER	23	-81.930	92.461	-22.595	1.00	78.00
50	ATOM	179	OG	SER	23	-81.564	93.718	-22.060	1.00	20.00
	ATOM	180	N	ASN	24	-80.865	94.517	-24.237	1.00	87.06
	ATOM	181	CA	ASN	24	-80.525	95.742	-24.933	1.00	88.17
	ATOM	182	C	ASN	24	-80.180	95.405	-26.392	1.00	88.45
	ATOM	183	O	ASN	24	-79.773	96.283	-27.132	1.00	84.89
55	ATOM	184	CB	ASN	24	-79.304	96.411	-24.343	0.00	86.84
	ATOM	185	CG	ASN	24	-79.428	96.644	-22.881	0.00	85.55
	ATOM	186	OD1	ASN	24	-80.536	96.786	-22.369	1.00	20.00
	ATOM	187	ND2	ASN	24	-78.304	96.693	-22.179	1.00	20.00
	ATOM	188	N	THR	25	-80.342	94.116	-26.792	1.00	89.49
60	ATOM	189	CA	THR	25	-79.860	93.730	-28.135	1.00	91.86
	ATOM	190	C	THR	25	-80.862	93.518	-29.320	1.00	95.75
	ATOM	191	O	THR	25	-80.588	92.760	-30.266	1.00	96.59
	ATOM	192	CB	THR	25	-78.942	92.539	-27.960	0.00	91.23
	ATOM	193	OG1	THR	25	-77.911	92.877	-27.030	1.00	20.00
65	ATOM	194	CG2	THR	25	-78.342	92.147	-29.292	1.00	20.00
	ATOM	195	N	PRO	26	-82.032	94.193	-29.248	1.00	98.41

	ATOM	196	CA	PRO	26	-83.068	94.281	-30.417	1.00	96.39
	ATOM	197	C	PRO	26	-82.814	95.192	-31.785	1.00	96.88
	ATOM	198	O	PRO	26	-83.158	96.376	-31.724	1.00	94.52
	ATOM	199	CB	PRO	26	-84.297	94.734	-29.653	0.00	95.29
5	ATOM	200	CG	PRO	26	-84.210	94.008	-28.343	0.00	92.34
	ATOM	201	CD	PRO	26	-82.772	93.675	-28.099	0.00	89.50
	ATOM	202	N	PRO	27	-82.213	94.701	-32.982	1.00	97.72
	ATOM	203	CA	PRO	27	-82.052	95.496	-34.330	1.00	98.94
	ATOM	204	C	PRO	27	-83.283	96.049	-35.168	1.00	100.71
10	ATOM	205	O	PRO	27	-84.403	95.618	-34.904	1.00	100.56
	ATOM	206	CB	PRO	27	-81.124	94.592	-35.110	0.00	97.33
	ATOM	207	CG	PRO	27	-80.205	94.081	-34.048	0.00	94.51
	ATOM	208	CD	PRO	27	-80.906	94.117	-32.739	0.00	92.58
	ATOM	209	N	LEU	28	-83.145	96.997	-36.177	1.00	15.00
15	ATOM	210	CA	LEU	28	-84.375	97.282	-36.988	1.00	15.00
	ATOM	211	C	LEU	28	-84.753	95.855	-37.376	1.00	15.00
	ATOM	212	O	LEU	28	-84.320	94.906	-36.721	1.00	15.00
	ATOM	213	CB	LEU	28	-84.124	98.182	-38.247	1.00	15.00
	ATOM	214	CG	LEU	28	-84.867	99.552	-38.338	1.00	15.00
20	ATOM	215	CD1	LEU	28	-85.868	99.526	-39.469	1.00	15.00
	ATOM	216	CD2	LEU	28	-85.547	99.887	-37.012	1.00	15.00
	ATOM	217	N	THR	29	-85.521	95.715	-38.443	1.00	15.00
	ATOM	218	CA	THR	29	-85.845	94.378	-38.924	1.00	15.00
	ATOM	219	C	THR	29	-85.688	93.353	-37.806	1.00	15.00
25	ATOM	220	O	THR	29	-85.492	92.176	-38.060	1.00	15.00
	ATOM	221	CB	THR	29	-84.961	94.028	-40.128	1.00	15.00
	ATOM	222	OG1	THR	29	-85.706	94.228	-41.325	1.00	20.00
	ATOM	223	CG2	THR	29	-84.502	92.579	-40.029	1.00	20.00
	ATOM	224	N	CYS	30	-85.794	93.855	-36.564	1.00	15.00
30	ATOM	225	CA	CYS	30	-85.616	92.973	-35.427	1.00	15.00
	ATOM	226	C	CYS	30	-86.621	93.301	-34.331	1.00	15.00
	ATOM	227	O	CYS	30	-87.018	92.421	-33.585	1.00	15.00
	ATOM	228	CB	CYS	30	-84.172	93.009	-34.948	1.00	15.00
	ATOM	229	SG	CYS	30	-82.984	92.154	-36.030	1.00	20.00
35	ATOM	230	N	GLN	31	-87.030	94.568	-34.256	1.00	15.00
	ATOM	231	CA	GLN	31	-88.017	94.944	-33.272	1.00	15.00
	ATOM	232	C	GLN	31	-89.312	94.148	-33.449	1.00	15.00
	ATOM	233	O	GLN	31	-89.853	93.572	-32.499	1.00	15.00
	ATOM	234	CB	GLN	31	-88.295	96.424	-33.328	1.00	15.00
40	ATOM	235	CG	GLN	31	-87.301	97.204	-32.478	1.00	20.00
	ATOM	236	CD	GLN	31	-87.610	98.686	-32.423	1.00	20.00
	ATOM	237	OE1	GLN	31	-88.729	99.108	-32.715	1.00	20.00
	ATOM	238	NE2	GLN	31	-86.770	99.655	-32.077	1.00	20.00
	ATOM	239	N	ARG	32	-89.795	94.113	-34.659	1.00	15.00
45	ATOM	240	CA	ARG	32	-90.985	93.365	-35.008	1.00	15.00
	ATOM	241	C	ARG	32	-90.848	91.886	-34.663	1.00	15.00
	ATOM	242	O	ARG	32	-91.796	91.250	-34.169	1.00	15.00
	ATOM	243	CB	ARG	32	-91.325	93.593	-36.496	1.00	15.00
	ATOM	244	CG	ARG	32	-91.631	95.047	-36.870	1.00	15.00
50	ATOM	245	CD	ARG	32	-92.030	95.147	-38.339	1.00	15.00
	ATOM	246	NE	ARG	32	-92.262	96.521	-38.769	1.00	20.00
	ATOM	247	CZ	ARG	32	-92.677	96.859	-39.984	1.00	20.00
	ATOM	248	NH1	ARG	32	-92.907	95.918	-40.891	1.00	20.00
	ATOM	249	NH2	ARG	32	-92.865	98.135	-40.292	1.00	20.00
55	ATOM	250	N	TYR	33	-89.648	91.370	-34.916	1.00	15.00
	ATOM	251	CA	TYR	33	-89.345	89.979	-34.626	1.00	15.00
	ATOM	252	C	TYR	33	-89.362	89.674	-33.102	1.00	15.00
	ATOM	253	O	TYR	33	-90.005	88.721	-32.681	1.00	15.00
	ATOM	254	CB	TYR	33	-88.021	89.652	-35.240	1.00	15.00
60	ATOM	255	CG	TYR	33	-87.988	89.919	-36.734	1.00	20.00
	ATOM	256	CD1	TYR	33	-87.579	91.138	-37.254	1.00	20.00
	ATOM	257	CD2	TYR	33	-88.349	88.915	-37.623	1.00	20.00
	ATOM	258	CE1	TYR	33	-87.529	91.352	-38.617	1.00	20.00
	ATOM	259	CE2	TYR	33	-88.300	89.119	-38.989	1.00	20.00
65	ATOM	260	CZ	TYR	33	-87.890	90.338	-39.481	1.00	20.00
	ATOM	261	OH	TYR	33	-87.839	90.546	-40.841	1.00	20.00

	ATOM	262	N	CYS	34	-88.662	90.482	-32.288	1.00	15.00
	ATOM	263	CA	CYS	34	-88.640	90.298	-30.845	1.00	15.00
	ATOM	264	C	CYS	34	-90.093	90.256	-30.311	1.00	15.00
	ATOM	265	O	CYS	34	-90.398	89.500	-29.388	1.00	15.00
5	ATOM	266	CB	CYS	34	-87.804	91.380	-30.172	1.00	15.00
	ATOM	267	SG	CYS	34	-86.025	91.318	-30.550	1.00	20.00
	ATOM	268	N	GLN	35	-90.959	91.067	-30.912	1.00	15.00
	ATOM	269	CA	GLN	35	-92.400	91.195	-30.611	1.00	15.00
	ATOM	270	CB	GLN	35	-92.927	92.414	-31.370	1.00	15.00
10	ATOM	271	CG	GLN	35	-94.239	92.978	-30.889	1.00	15.00
	ATOM	272	CD	GLN	35	-94.225	93.168	-29.401	1.00	15.00
	ATOM	273	OE1	GLN	35	-93.330	93.817	-28.865	1.00	15.00
	ATOM	274	NE2	GLN	35	-95.113	92.712	-28.540	1.00	15.00
	ATOM	275	C	GLN	35	-93.205	89.974	-31.045	1.00	15.00
15	ATOM	276	O	GLN	35	-94.029	89.457	-30.285	1.00	15.00
	ATOM	277	N	ALA	36	-92.927	89.534	-32.267	1.00	15.00
	ATOM	278	CA	ALA	36	-93.535	88.342	-32.864	1.00	15.00
	ATOM	279	C	ALA	36	-93.327	87.151	-31.911	1.00	15.00
	ATOM	280	O	ALA	36	-94.280	86.435	-31.595	1.00	15.00
20	ATOM	281	CB	ALA	36	-92.942	88.047	-34.219	1.00	15.00
	ATOM	282	N	SER	37	-92.053	86.955	-31.492	1.00	15.00
	ATOM	283	CA	SER	37	-91.761	85.801	-30.609	1.00	15.00
	ATOM	284	C	SER	37	-92.380	86.039	-29.251	1.00	15.00
	ATOM	285	O	SER	37	-92.916	85.109	-28.626	1.00	15.00
25	ATOM	286	CB	SER	37	-90.253	85.510	-30.545	1.00	15.00
	ATOM	287	OG	SER	37	-89.503	86.681	-30.277	1.00	20.00
	ATOM	288	N	VAL	38	-92.306	87.290	-28.785	1.00	15.00
	ATOM	289	CA	VAL	38	-92.897	87.655	-27.528	1.00	15.00
	ATOM	290	C	VAL	38	-94.380	87.268	-27.466	1.00	15.00
30	ATOM	291	O	VAL	38	-94.846	86.602	-26.522	1.00	15.00
	ATOM	292	CB	VAL	38	-92.769	89.189	-27.293	1.00	15.00
	ATOM	293	CG1	VAL	38	-93.640	89.645	-26.122	1.00	20.00
	ATOM	294	CG2	VAL	38	-91.313	89.563	-27.030	1.00	20.00
	ATOM	295	N	THR	39	-95.124	87.731	-28.485	1.00	15.00
35	ATOM	296	CA	THR	39	-96.560	87.468	-28.556	1.00	15.00
	ATOM	297	C	THR	39	-96.849	86.015	-28.210	1.00	15.00
	ATOM	298	O	THR	39	-98.012	85.612	-28.115	1.00	15.00
	ATOM	299	CB	THR	39	-97.073	87.777	-29.971	1.00	15.00
	ATOM	300	OG1	THR	39	-96.899	89.176	-30.251	1.00	20.00
40	ATOM	301	CG2	THR	39	-98.531	87.397	-30.081	1.00	20.00
	END					-118.331	85.231	-119.150	0.00	0.00

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	ATOM	1	C	CYS	1	10.780	107.980	37.608 1.00101.54
5	ATOM	2	O	CYS	1	10.355	106.862	37.325 1.00102.94
	ATOM	3	CB	CYS	1	10.689	109.854	35.953 1.00 96.91
	ATOM	4	SG	CYS	1	11.113	111.582	36.339 1.00 92.39
	ATOM	5	N	CYS	1	12.253	107.967	35.619 1.00 93.44
	ATOM	6	CA	CYS	1	11.595	108.804	36.629 1.00 98.53
10	ATOM	7	N	SER	2	10.584	108.616	38.768 1.00104.82
	ATOM	8	CA	SER	2	9.771	108.207	39.905 1.00107.58
	ATOM	9	CB	SER	2	9.995	106.748	40.341 1.00109.37
	ATOM	10	OG	SER	2	8.782	106.129	40.715 1.00110.15
	ATOM	11	C	SER	2	10.051	109.213	41.053 1.00108.57
15	ATOM	12	O	SER	2	9.692	110.384	40.910 1.00109.82
	ATOM	13	N	GLN	3	10.665	108.806	42.165 1.00108.52
	ATOM	14	CA	GLN	3	10.831	109.704	43.324 1.00109.56
	ATOM	15	CB	GLN	3	12.072	110.608	43.199 1.00108.23
	ATOM	16	CG	GLN	3	11.932	111.788	42.262 1.00109.80
20	ATOM	17	CD	GLN	3	12.683	111.497	40.995 1.00110.35
	ATOM	18	OE1	GLN	3	13.883	111.248	40.997 1.00110.92
	ATOM	19	NE2	GLN	3	12.156	111.479	39.781 1.00110.59
	ATOM	20	C	GLN	3	9.528	110.507	43.566 1.00110.79
	ATOM	21	O	GLN	3	9.526	111.742	43.608 1.00114.50
25	ATOM	22	N	ASN	4	8.422	109.762	43.726 1.00109.87
	ATOM	23	CA	ASN	4	7.029	110.198	44.011 1.00107.78
	ATOM	24	CB	ASN	4	6.887	110.535	45.489 1.00109.08
	ATOM	25	CG	ASN	4	7.952	111.491	46.016 1.00109.99
	ATOM	26	OD1	ASN	4	9.124	111.133	46.139 1.00109.73
30	ATOM	27	ND2	ASN	4	7.531	112.717	46.336 1.00109.43
	ATOM	28	C	ASN	4	6.518	111.329	43.113 1.00105.15
	ATOM	29	O	ASN	4	6.223	112.417	43.581 1.00103.42
	ATOM	30	N	GLU	5	6.433	111.064	41.785 1.00101.11
	ATOM	31	CA	GLU	5	5.952	112.085	40.794 1.00 95.31
35	ATOM	32	CB	GLU	5	7.084	113.040	40.417 1.00 92.47
	ATOM	33	CG	GLU	5	8.133	113.268	41.494 1.00 86.82
	ATOM	34	CD	GLU	5	9.519	113.154	40.891 1.00 85.55
	ATOM	35	OE1	GLU	5	9.733	112.225	40.093 1.00 85.72
	ATOM	36	OE2	GLU	5	10.366	113.996	41.223 1.00 81.01
40	ATOM	37	C	GLU	5	5.329	111.536	39.485 1.00 92.66
	ATOM	38	O	GLU	5	4.819	110.431	39.431 1.00 94.43
	ATOM	39	N	TYR	6	5.399	112.374	38.451 1.00 85.19
	ATOM	40	CA	TYR	6	4.853	112.065	37.127 1.00 78.14
	ATOM	41	CB	TYR	6	3.400	112.492	37.031 1.00 76.59
45	ATOM	42	CG	TYR	6	3.185	113.963	36.857 1.00 73.94
	ATOM	43	CD1	TYR	6	2.880	114.464	35.583 1.00 73.49
	ATOM	44	CE1	TYR	6	2.654	115.813	35.392 1.00 73.13
	ATOM	45	CD2	TYR	6	3.266	114.865	37.922 1.00 72.45
	ATOM	46	CE2	TYR	6	3.040	116.236	37.731 1.00 72.38
50	ATOM	47	CZ	TYR	6	2.731	116.693	36.473 1.00 74.15
	ATOM	48	OH	TYR	6	2.509	118.035	36.267 1.00 73.32
	ATOM	49	C	TYR	6	5.694	112.775	36.090 1.00 75.43
	ATOM	50	O	TYR	6	6.406	113.744	36.416 1.00 76.93
	ATOM	51	N	PHE	7	5.628	112.320	34.842 1.00 69.11
55	ATOM	52	CA	PHE	7	6.385	112.973	33.780 1.00 62.54
	ATOM	53	CB	PHE	7	7.067	111.912	32.911 1.00 61.14
	ATOM	54	CG	PHE	7	7.780	112.516	31.731 1.00 61.11
	ATOM	55	CD1	PHE	7	9.001	113.149	31.928 1.00 59.90
	ATOM	56	CD2	PHE	7	7.235	112.489	30.448 1.00 63.23
60	ATOM	57	CE1	PHE	7	9.671	113.751	30.862 1.00 61.45
	ATOM	58	CE2	PHE	7	7.892	113.088	29.366 1.00 61.15
	ATOM	59	CZ	PHE	7	9.110	113.720	29.572 1.00 61.89
	ATOM	60	C	PHE	7	5.483	113.884	32.933 1.00 60.79
	ATOM	61	O	PHE	7	4.820	113.412	32.005 1.00 63.10
65	ATOM	62	N	ASP	8	5.456	115.183	33.248 1.00 57.06
	ATOM	63	CA	ASP	8	4.633	116.145	32.506 1.00 52.20

	ATOM	64	CB	ASP	8	4.607	117.504	33.200	1.00	46.52
	ATOM	65	CG	ASP	8	3.573	118.476	32.651	1.00	47.00
	ATOM	66	OD1	ASP	8	3.325	118.430	31.426	1.00	49.00
	ATOM	67	OD2	ASP	8	3.031	119.276	33.441	1.00	48.43
5	ATOM	68	C	ASP	8	5.176	116.300	31.079	1.00	53.02
	ATOM	69	O	ASP	8	6.258	116.860	30.872	1.00	53.25
	ATOM	70	N	SER	9	4.425	115.795	30.100	1.00	55.34
	ATOM	71	CA	SER	9	4.825	115.882	28.690	1.00	54.13
	ATOM	72	CB	SER	9	3.885	115.057	27.805	1.00	54.65
10	ATOM	73	OG	SER	9	3.920	113.690	28.174	1.00	54.09
	ATOM	74	C	SER	9	4.835	117.325	28.180	1.00	54.96
	ATOM	75	O	SER	9	5.577	117.656	27.240	1.00	54.11
	ATOM	76	N	LEU	10	4.030	118.192	28.786	1.00	49.64
	ATOM	77	CA	LEU	10	4.029	119.574	28.353	1.00	46.11
15	ATOM	78	CB	LEU	10	2.952	120.384	29.060	1.00	40.39
	ATOM	79	CG	LEU	10	2.725	121.786	28.538	1.00	34.58
	ATOM	80	CD1	LEU	10	2.141	121.740	27.124	1.00	33.04
	ATOM	81	CD2	LEU	10	1.818	122.580	29.481	1.00	33.82
	ATOM	82	C	LEU	10	5.425	120.139	28.619	1.00	49.82
20	ATOM	83	O	LEU	10	6.004	120.795	27.769	1.00	50.94
	ATOM	84	N	LEU	11	5.958	119.881	29.813	1.00	51.20
	ATOM	85	CA	LEU	11	7.278	120.391	30.178	1.00	51.39
	ATOM	86	CB	LEU	11	7.363	120.606	31.693	1.00	47.82
	ATOM	87	CG	LEU	11	6.314	121.512	32.355	1.00	46.28
25	ATOM	88	CD1	LEU	11	6.599	121.646	33.847	1.00	49.44
	ATOM	89	CD2	LEU	11	6.280	122.885	31.699	1.00	44.53
	ATOM	90	C	LEU	11	8.454	119.516	29.729	1.00	53.87
	ATOM	91	O	LEU	11	9.573	119.997	29.600	1.00	50.27
	ATOM	92	N	HIS	12	8.201	118.223	29.504	1.00	58.57
30	ATOM	93	CA	HIS	12	9.229	117.278	29.119	1.00	64.33
	ATOM	94	CB	HIS	12	9.997	117.746	27.884	1.00	65.68
	ATOM	95	CG	HIS	12	9.125	117.781	26.602	1.00	69.20
	ATOM	96	CD2	HIS	12	8.065	117.024	26.227	1.00	69.34
	ATOM	97	ND1	HIS	12	9.315	118.699	25.592	1.00	71.64
35	ATOM	98	CE1	HIS	12	8.402	118.513	24.650	1.00	72.29
	ATOM	99	NE2	HIS	12	7.632	117.501	25.012	1.00	70.37
	ATOM	100	C	HIS	12	10.242	117.116	30.243	1.00	66.69
	ATOM	101	O	HIS	12	11.458	117.066	30.030	1.00	67.11
	ATOM	102	N	ALA	13	9.711	117.057	31.461	1.00	69.57
40	ATOM	103	CA	ALA	13	10.521	116.939	32.666	1.00	71.62
	ATOM	104	CB	ALA	13	10.972	118.295	33.155	1.00	69.66
	ATOM	105	C	ALA	13	9.701	116.224	33.736	1.00	73.99
	ATOM	106	O	ALA	13	8.568	115.802	33.475	1.00	75.45
	ATOM	107	N	CYS	14	10.254	116.102	34.941	1.00	75.75
45	ATOM	108	CA	CYS	14	9.547	115.409	36.011	1.00	77.01
	ATOM	109	C	CYS	14	9.077	116.302	37.140	1.00	75.86
	ATOM	110	O	CYS	14	9.863	116.960	37.827	1.00	73.08
	ATOM	111	CB	CYS	14	10.413	114.261	36.534	1.00	80.96
	ATOM	112	SG	CYS	14	10.008	112.632	35.834	1.00	86.58
50	ATOM	113	N	ILE	15	7.737	116.292	37.313	1.00	75.89
	ATOM	114	CA	ILE	15	7.102	117.120	38.325	1.00	77.02
	ATOM	115	CB	ILE	15	5.973	117.982	37.712	1.00	76.68
	ATOM	116	CG2	ILE	15	5.305	118.830	38.790	1.00	76.37
	ATOM	117	CG1	ILE	15	6.508	118.881	36.583	1.00	76.28
55	ATOM	118	CD1	ILE	15	7.490	119.941	37.035	1.00	74.77
	ATOM	119	C	ILE	15	6.541	116.314	39.520	1.00	78.19
	ATOM	120	O	ILE	15	5.894	115.287	39.377	1.00	77.54
	ATOM	121	N	PRO	16	6.830	116.889	40.718	1.00	80.09
	ATOM	122	CD	PRO	16	8.192	117.406	40.987	1.00	79.64
60	ATOM	123	CA	PRO	16	6.298	116.299	41.983	1.00	82.16
	ATOM	124	CB	PRO	16	6.863	117.233	43.039	1.00	80.00
	ATOM	125	CG	PRO	16	8.208	117.552	42.482	1.00	80.17
	ATOM	126	C	PRO	16	4.822	116.028	41.958	1.00	84.28
	ATOM	127	O	PRO	16	4.050	116.946	41.688	1.00	84.12
65	ATOM	128	N	CYS	17	4.384	114.792	42.221	1.00	87.37
	ATOM	129	CA	CYS	17	2.961	114.495	42.203	1.00	88.33

	ATOM	130	CB	CYS	17	2.703	113.091	42.741	1.00	86.90
	ATOM	131	SG	CYS	17	3.026	111.756	41.551	1.00	84.56
	ATOM	132	C	CYS	17	2.160	115.521	43.014	1.00	89.64
	ATOM	133	O	CYS	17	1.060	115.934	42.611	1.00	89.77
5	ATOM	134	N	GLN	18	2.722	115.919	44.156	1.00	91.08
	ATOM	135	CA	GLN	18	2.109	116.900	45.003	1.00	91.16
	ATOM	136	CB	GLN	18	3.119	117.562	45.942	1.00	92.68
	ATOM	137	CG	GLN	18	4.123	116.574	46.524	1.00	98.31
	ATOM	138	CD	GLN	18	5.432	117.210	47.009	1.00	101.37
10	ATOM	139	OE1	GLN	18	6.240	117.672	46.212	1.00	102.74
	ATOM	140	NE2	GLN	18	5.817	117.330	48.266	1.00	103.76
	ATOM	141	C	GLN	18	1.456	117.968	44.121	1.00	89.18
	ATOM	142	O	GLN	18	0.259	118.245	44.222	1.00	89.64
	ATOM	143	N	LEU	19	2.294	118.554	43.265	1.00	87.56
15	ATOM	144	CA	LEU	19	1.911	119.626	42.363	1.00	85.33
	ATOM	145	CB	LEU	19	3.026	119.903	41.396	1.00	79.98
	ATOM	146	CG	LEU	19	3.827	121.142	41.735	1.00	77.91
	ATOM	147	CD1	LEU	19	5.140	121.158	40.970	1.00	78.13
	ATOM	148	CD2	LEU	19	3.026	122.400	41.452	1.00	72.50
20	ATOM	149	C	LEU	19	0.621	119.393	41.596	1.00	86.05
	ATOM	150	O	LEU	19	-0.219	120.302	41.505	1.00	88.06
	ATOM	151	N	ARG	20	0.438	118.212	41.058	1.00	85.11
	ATOM	152	CA	ARG	20	-0.772	117.955	40.266	1.00	83.33
	ATOM	153	CB	ARG	20	-0.522	116.819	39.278	1.00	78.93
25	ATOM	154	CG	ARG	20	-0.305	117.267	37.836	1.00	73.68
	ATOM	155	CD	ARG	20	-1.563	117.866	37.233	1.00	67.23
	ATOM	156	NE	ARG	20	-2.079	117.056	36.129	1.00	63.84
	ATOM	157	CZ	ARG	20	-1.336	116.619	35.109	1.00	64.39
	ATOM	158	NH1	ARG	20	-0.042	116.921	35.067	1.00	66.99
30	ATOM	159	NH2	ARG	20	-1.889	115.896	34.149	1.00	60.63
	ATOM	160	C	ARG	20	-2.009	117.665	41.117	1.00	84.21
	ATOM	161	O	ARG	20	-2.978	117.108	40.602	1.00	86.28
	ATOM	162	N	CYS	21	-1.989	118.029	42.391	1.00	85.02
	ATOM	163	CA	CYS	21	-3.173	117.792	43.209	1.00	83.51
35	ATOM	164	CB	CYS	21	-2.775	117.014	44.471	1.00	81.05
	ATOM	165	SG	CYS	21	-1.973	115.412	44.147	1.00	73.08
	ATOM	166	C	CYS	21	-3.942	119.095	43.549	1.00	84.26
	ATOM	167	O	CYS	21	-3.948	119.498	44.703	1.00	84.48
	ATOM	168	N	SER	22	-4.586	119.779	42.538	1.00	84.61
40	ATOM	169	CA	SER	22	-5.380	121.038	42.693	1.00	85.07
	ATOM	170	C	SER	22	-6.855	120.767	42.436	1.00	85.65
	ATOM	171	O	SER	22	-7.632	121.649	42.079	1.00	86.77
	ATOM	172	CB	SER	22	-4.917	122.126	41.715	1.00	20.00
	ATOM	173	OG	SER	22	-5.581	122.002	40.472	1.00	20.00
45	ATOM	174	N	SER	23	-7.177	119.520	42.647	1.00	84.19
	ATOM	175	CA	SER	23	-8.540	119.056	42.523	1.00	82.44
	ATOM	176	C	SER	23	-8.825	118.322	41.209	1.00	84.32
	ATOM	177	O	SER	23	-8.508	117.135	41.113	1.00	81.24
	ATOM	178	CB	SER	23	-9.474	120.245	42.747	1.00	78.00
50	ATOM	179	OG	SER	23	-10.618	120.168	41.920	1.00	20.00
	ATOM	180	N	ASN	24	-9.392	118.954	40.221	1.00	87.06
	ATOM	181	CA	ASN	24	-9.605	118.159	39.028	1.00	88.17
	ATOM	182	C	ASN	24	-8.243	117.699	38.488	1.00	88.45
	ATOM	183	O	ASN	24	-8.187	117.093	37.432	1.00	84.89
55	ATOM	184	CB	ASN	24	-10.261	118.957	37.923	0.00	86.84
	ATOM	185	CG	ASN	24	-11.520	119.613	38.360	0.00	85.55
	ATOM	186	OD1	ASN	24	-12.187	119.128	39.271	1.00	20.00
	ATOM	187	ND2	ASN	24	-11.877	120.721	37.724	1.00	20.00
	ATOM	188	N	THR	25	-7.142	118.000	39.226	1.00	89.49
60	ATOM	189	CA	THR	25	-5.810	117.720	38.651	1.00	91.86
	ATOM	190	C	THR	25	-4.967	116.490	39.131	1.00	95.75
	ATOM	191	O	THR	25	-3.727	116.491	39.057	1.00	96.59
	ATOM	192	CB	THR	25	-5.011	119.004	38.716	0.00	91.23
	ATOM	193	OG1	THR	25	-5.738	120.042	38.057	1.00	20.00
65	ATOM	194	CG2	THR	25	-3.661	118.808	38.060	1.00	20.00
	ATOM	195	N	PRO	26	-5.662	115.444	39.635	1.00	98.41

	ATOM	196	CA	PRO	26	-5.032	114.052	39.971	1.00	96.39
	ATOM	197	C	PRO	26	-4.558	112.966	38.804	1.00	96.88
	ATOM	198	O	PRO	26	-5.427	112.193	38.392	1.00	94.52
	ATOM	199	CB	PRO	26	-6.108	113.482	40.874	0.00	95.29
5	ATOM	200	CG	PRO	26	-6.598	114.661	41.661	0.00	92.34
	ATOM	201	CD	PRO	26	-6.313	115.894	40.864	0.00	89.50
	ATOM	202	N	PRO	27	-3.246	112.900	38.249	1.00	97.72
	ATOM	203	CA	PRO	27	-2.728	111.823	37.225	1.00	98.94
	ATOM	204	C	PRO	27	-2.678	110.268	37.546	1.00	100.71
10	ATOM	205	O	PRO	27	-2.793	109.915	38.718	1.00	100.56
	ATOM	206	CB	PRO	27	-1.402	112.415	36.804	0.00	97.33
	ATOM	207	CG	PRO	27	-1.700	113.878	36.764	0.00	94.51
	ATOM	208	CD	PRO	27	-2.820	114.172	37.695	0.00	92.58
	ATOM	209	N	LEU	28	-2.515	109.299	36.561	1.00	15.00
15	ATOM	210	CA	LEU	28	-2.312	107.893	37.045	1.00	15.00
	ATOM	211	C	LEU	28	-1.169	108.132	38.030	1.00	15.00
	ATOM	212	O	LEU	28	-0.968	109.263	38.472	1.00	15.00
	ATOM	213	CB	LEU	28	-1.912	106.875	35.923	1.00	15.00
	ATOM	214	CG	LEU	28	-2.859	105.663	35.650	1.00	15.00
20	ATOM	215	CD1	LEU	28	-2.177	104.372	36.040	1.00	15.00
	ATOM	216	CD2	LEU	28	-4.180	105.832	36.397	1.00	15.00
	ATOM	217	N	THR	29	-0.421	107.085	38.332	1.00	15.00
	ATOM	218	CA	THR	29	0.743	107.260	39.193	1.00	15.00
	ATOM	219	C	THR	29	0.601	108.521	40.037	1.00	15.00
25	ATOM	220	O	THR	29	1.584	109.083	40.492	1.00	15.00
	ATOM	221	CB	THR	29	2.019	107.301	38.342	1.00	15.00
	ATOM	222	OG1	THR	29	2.649	106.024	38.388	1.00	20.00
	ATOM	223	CG2	THR	29	2.961	108.373	38.875	1.00	20.00
	ATOM	224	N	CYS	30	-0.665	108.927	40.233	1.00	15.00
30	ATOM	225	CA	CYS	30	-0.911	110.142	40.985	1.00	15.00
	ATOM	226	C	CYS	30	-2.114	109.970	41.903	1.00	15.00
	ATOM	227	O	CYS	30	-2.170	110.587	42.955	1.00	15.00
	ATOM	228	CB	CYS	30	-1.036	111.332	40.046	1.00	15.00
	ATOM	229	SG	CYS	30	0.529	111.891	39.303	1.00	20.00
35	ATOM	230	N	GLN	31	-3.059	109.123	41.496	1.00	15.00
	ATOM	231	CA	GLN	31	-4.207	108.874	42.336	1.00	15.00
	ATOM	232	C	GLN	31	-3.789	108.333	43.705	1.00	15.00
	ATOM	233	O	GLN	31	-4.219	108.823	44.754	1.00	15.00
	ATOM	234	CB	GLN	31	-5.169	107.927	41.665	1.00	15.00
40	ATOM	235	CG	GLN	31	-6.129	108.676	40.750	1.00	20.00
	ATOM	236	CD	GLN	31	-7.181	107.773	40.137	1.00	20.00
	ATOM	237	OE1	GLN	31	-7.433	106.674	40.629	1.00	20.00
	ATOM	238	NE2	GLN	31	-7.916	108.036	39.062	1.00	20.00
	ATOM	239	N	ARG	32	-2.951	107.334	43.688	1.00	15.00
45	ATOM	240	CA	ARG	32	-2.419	106.735	44.894	1.00	15.00
	ATOM	241	C	ARG	32	-1.697	107.759	45.765	1.00	15.00
	ATOM	242	O	ARG	32	-1.820	107.752	47.002	1.00	15.00
	ATOM	243	CB	ARG	32	-1.519	105.537	44.527	1.00	15.00
	ATOM	244	CG	ARG	32	-2.233	104.400	43.787	1.00	15.00
50	ATOM	245	CD	ARG	32	-1.275	103.238	43.545	1.00	15.00
	ATOM	246	NE	ARG	32	-1.883	102.157	42.778	1.00	20.00
	ATOM	247	CZ	ARG	32	-1.270	101.013	42.492	1.00	20.00
	ATOM	248	NH1	ARG	32	-0.029	100.804	42.913	1.00	20.00
	ATOM	249	NH2	ARG	32	-1.898	100.080	41.790	1.00	20.00
55	ATOM	250	N	TYR	33	-0.965	108.640	45.087	1.00	15.00
	ATOM	251	CA	TYR	33	-0.230	109.694	45.765	1.00	15.00
	ATOM	252	C	TYR	33	-1.169	110.722	46.456	1.00	15.00
	ATOM	253	O	TYR	33	-0.980	111.026	47.626	1.00	15.00
	ATOM	254	CB	TYR	33	0.670	110.351	44.768	1.00	15.00
60	ATOM	255	CG	TYR	33	1.614	109.369	44.098	1.00	20.00
	ATOM	256	CD1	TYR	33	1.287	108.724	42.915	1.00	20.00
	ATOM	257	CD2	TYR	33	2.860	109.117	44.658	1.00	20.00
	ATOM	258	CE1	TYR	33	2.171	107.856	42.306	1.00	20.00
	ATOM	259	CE2	TYR	33	3.753	108.251	44.056	1.00	20.00
65	ATOM	260	CZ	TYR	33	3.404	107.624	42.880	1.00	20.00
	ATOM	261	OH	TYR	33	4.290	106.761	42.275	1.00	20.00

	ATOM	262	N	CYS	34	-2.173	111.246	45.731	1.00	15.00
	ATOM	263	CA	CYS	34	-3.123	112.193	46.296	1.00	15.00
	ATOM	264	C	CYS	34	-3.748	111.589	47.578	1.00	15.00
	ATOM	265	O	CYS	34	-4.001	112.306	48.547	1.00	15.00
5	ATOM	266	CB	CYS	34	-4.175	112.586	45.265	1.00	15.00
	ATOM	267	SG	CYS	34	-3.541	113.542	43.852	1.00	20.00
	ATOM	268	N	GLN	35	-3.977	110.279	47.554	1.00	15.00
	ATOM	269	CA	GLN	35	-4.537	109.463	48.652	1.00	15.00
	ATOM	270	CB	GLN	35	-4.852	108.077	48.087	1.00	15.00
10	ATOM	271	CG	GLN	35	-5.804	107.233	48.895	1.00	15.00
	ATOM	272	CD	GLN	35	-7.030	108.015	49.262	1.00	15.00
	ATOM	273	OE1	GLN	35	-7.691	108.581	48.394	1.00	15.00
	ATOM	274	NE2	GLN	35	-7.532	108.171	50.470	1.00	15.00
	ATOM	275	C	GLN	35	-3.568	109.298	49.819	1.00	15.00
15	ATOM	276	O	GLN	35	-3.944	109.468	50.982	1.00	15.00
	ATOM	277	N	ALA	36	-2.326	108.984	49.466	1.00	15.00
	ATOM	278	CA	ALA	36	-1.221	108.836	50.415	1.00	15.00
	ATOM	279	C	ALA	36	-1.124	110.116	51.265	1.00	15.00
	ATOM	280	O	ALA	36	-1.064	110.042	52.495	1.00	15.00
20	ATOM	281	CB	ALA	36	0.082	108.574	49.701	1.00	15.00
	ATOM	282	N	SER	37	-1.086	111.277	50.568	1.00	15.00
	ATOM	283	CA	SER	37	-0.946	112.552	51.310	1.00	15.00
	ATOM	284	C	SER	37	-2.218	112.827	52.078	1.00	15.00
	ATOM	285	O	SER	37	-2.178	113.308	53.224	1.00	15.00
25	ATOM	286	CB	SER	37	-0.541	113.704	50.377	1.00	15.00
	ATOM	287	OG	SER	37	-1.364	113.763	49.226	1.00	20.00
	ATOM	288	N	VAL	38	-3.359	112.526	51.450	1.00	15.00
	ATOM	289	CA	VAL	38	-4.632	112.696	52.091	1.00	15.00
	ATOM	290	C	VAL	38	-4.689	111.970	53.440	1.00	15.00
30	ATOM	291	O	VAL	38	-5.044	112.550	54.483	1.00	15.00
	ATOM	292	CB	VAL	38	-5.775	112.155	51.181	1.00	15.00
	ATOM	293	CG1	VAL	38	-7.092	112.050	51.948	1.00	20.00
	ATOM	294	CG2	VAL	38	-5.956	113.060	49.965	1.00	20.00
	ATOM	295	N	THR	39	-4.361	110.668	53.391	1.00	15.00
35	ATOM	296	CA	THR	39	-4.391	109.833	54.590	1.00	15.00
	ATOM	297	C	THR	39	-3.762	110.569	55.763	1.00	15.00
	ATOM	298	O	THR	39	-3.777	110.076	56.894	1.00	15.00
	ATOM	299	CB	THR	39	-3.629	108.524	54.328	1.00	15.00
	ATOM	300	OG1	THR	39	-4.293	107.779	53.294	1.00	20.00
40	ATOM	301	CG2	THR	39	-3.558	107.711	55.599	1.00	20.00
	END					60.588	44.144	42.385	0.00	0.00

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	ATOM	2	CA	PRO	1	-8.020	46.256	-51.178	1.00	98.53
	ATOM	3	C	PRO	1	-6.555	46.072	-51.070	1.00	101.54
5	ATOM	4	O	PRO	1	-5.930	45.970	-52.135	1.00	102.94
	ATOM	5	CB	PRO	1	-8.539	45.860	-52.546	1.00	96.91
	ATOM	6	CG	PRO	1	-9.873	46.515	-52.522	1.00	20.00
	ATOM	7	CD	PRO	1	-9.859	47.625	-51.483	1.00	20.00
	ATOM	8	N	THR	2	-5.905	46.032	-49.963	1.00	104.82
10	ATOM	9	CA	THR	2	-4.537	45.770	-50.271	1.00	107.58
	ATOM	10	C	THR	2	-4.265	44.324	-49.892	1.00	108.57
	ATOM	11	O	THR	2	-5.209	43.534	-49.918	1.00	109.82
	ATOM	12	CB	THR	2	-3.620	46.808	-49.648	1.00	20.00
	ATOM	13	OG1	THR	2	-3.491	47.935	-50.543	1.00	20.00
15	ATOM	14	CG2	THR	2	-2.257	46.199	-49.364	1.00	20.00
	ATOM	15	N	PRO	3	-3.033	43.886	-49.541	1.00	108.52
	ATOM	16	CA	PRO	3	-2.885	42.414	-49.386	1.00	109.56
	ATOM	17	C	PRO	3	-3.890	41.669	-48.573	1.00	110.79
	ATOM	18	O	PRO	3	-4.106	42.022	-47.414	1.00	114.50
20	ATOM	19	CB	PRO	3	-1.454	42.244	-48.916	1.00	108.23
	ATOM	20	CG	PRO	3	-0.772	43.280	-49.743	1.00	109.80
	ATOM	21	CD	PRO	3	-1.780	44.345	-50.127	1.00	110.35
	ATOM	22	C	CYS	4	-5.700	38.681	-47.973	1.00	101.54
	ATOM	23	O	CYS	4	-6.384	38.107	-48.829	1.00	102.94
25	ATOM	24	CB	CYS	4	-6.582	41.024	-47.911	1.00	96.91
	ATOM	25	SG	CYS	4	-6.692	41.947	-46.345	1.00	92.39
	ATOM	26	N	CYS	4	-4.514	40.625	-49.070	1.00	93.44
	ATOM	27	CA	CYS	4	-5.378	40.158	-47.992	1.00	98.53
	ATOM	28	N	VAL	5	-5.180	38.135	-46.900	1.00	104.82
30	ATOM	29	CA	VAL	5	-5.247	36.733	-46.574	1.00	107.58
	ATOM	30	C	VAL	5	-6.704	36.214	-46.414	1.00	108.57
	ATOM	31	O	VAL	5	-7.556	36.917	-45.862	1.00	109.82
	ATOM	32	CB	VAL	5	-4.273	36.546	-45.400	1.00	109.37
	ATOM	33	CG1	VAL	5	-3.981	35.069	-45.182	1.00	20.00
35	ATOM	34	CG2	VAL	5	-2.989	37.327	-45.617	1.00	20.00
	ATOM	35	N	PRO	6	-6.965	34.989	-46.918	1.00	108.52
	ATOM	36	CA	PRO	6	-8.362	34.509	-46.818	1.00	109.56
	ATOM	37	C	PRO	6	-9.016	34.668	-45.438	1.00	110.79
	ATOM	38	O	PRO	6	-8.382	34.400	-44.427	1.00	114.50
40	ATOM	39	CB	PRO	6	-8.294	33.121	-47.438	1.00	108.23
	ATOM	40	CG	PRO	6	-7.340	33.303	-48.560	1.00	109.80
	ATOM	41	CD	PRO	6	-6.456	34.482	-48.214	1.00	110.35
	ATOM	42	N	ALA	7	-10.280	35.081	-45.384	1.00	109.87
	ATOM	43	CA	ALA	7	-10.984	35.347	-44.130	1.00	107.78
45	ATOM	44	C	ALA	7	-10.949	36.834	-43.729	1.00	105.15
	ATOM	45	O	ALA	7	-11.976	37.353	-43.326	1.00	103.42
	ATOM	46	CB	ALA	7	-10.414	34.495	-43.014	1.00	109.08
	ATOM	47	N	GLU	8	-9.807	37.582	-43.866	1.00	101.11
	ATOM	48	CA	GLU	8	-9.776	39.015	-43.364	1.00	95.31
50	ATOM	49	CB	GLU	8	-8.481	39.357	-42.674	1.00	92.47
	ATOM	50	CG	GLU	8	-7.561	38.174	-42.436	1.00	86.82
	ATOM	51	CD	GLU	8	-6.129	38.481	-42.839	1.00	85.55
	ATOM	52	OE1	GLU	8	-5.907	38.892	-44.002	1.00	85.72
	ATOM	53	OE2	GLU	8	-5.236	38.331	-41.993	1.00	81.01
55	ATOM	54	C	GLU	8	-10.041	40.135	-44.352	1.00	92.66
	ATOM	55	O	GLU	8	-9.979	39.950	-45.573	1.00	94.43
	ATOM	56	N	CYS	9	-10.331	41.309	-43.803	1.00	85.19
	ATOM	57	CA	CYS	9	-10.653	42.475	-44.611	1.00	78.14
	ATOM	58	C	CYS	9	-9.756	43.621	-44.190	1.00	75.43
60	ATOM	59	O	CYS	9	-9.528	43.843	-43.004	1.00	76.93
	ATOM	60	CB	CYS	9	-12.119	42.868	-44.439	1.00	76.59
	ATOM	61	SG	CYS	9	-12.543	43.520	-42.793	1.00	20.00
	ATOM	62	N	PHE	10	-9.257	44.334	-45.199	1.00	69.11
	ATOM	63	CA	PHE	10	-8.395	45.528	-45.019	1.00	62.54
65	ATOM	64	CB	PHE	10	-7.855	46.030	-46.413	1.00	61.14
	ATOM	65	CG	PHE	10	-6.898	47.199	-46.340	1.00	61.11

	ATOM	66	CD1	PHE	10	-5.623	46.996	-45.827	1.00	59.90
	ATOM	67	CD2	PHE	10	-7.271	48.485	-46.728	1.00	63.23
	ATOM	68	CE1	PHE	10	-4.729	48.061	-45.700	1.00	61.45
	ATOM	69	CE2	PHE	10	-6.388	49.566	-46.608	1.00	61.15
5	ATOM	70	CZ	PHE	10	-5.117	49.355	-46.093	1.00	61.89
	ATOM	71	C	PHE	10	-9.189	46.644	-44.318	1.00	60.79
	ATOM	72	O	PHE	10	-10.197	47.131	-44.840	1.00	63.10
	ATOM	73	N	ASP	11	-8.756	47.043	-43.115	1.00	57.06
	ATOM	74	CA	ASP	11	-9.432	48.138	-42.388	1.00	52.20
10	ATOM	75	CB	ASP	11	-9.352	47.921	-40.829	1.00	46.52
	ATOM	76	CG	ASP	11	-10.114	48.937	-39.959	1.00	47.00
	ATOM	77	OD1	ASP	11	-10.206	50.111	-40.382	1.00	49.00
	ATOM	78	OD2	ASP	11	-10.609	48.544	-38.884	1.00	48.43
	ATOM	79	C	ASP	11	-8.724	49.453	-42.773	1.00	53.02
15	ATOM	80	O	ASP	11	-7.534	49.587	-42.503	1.00	53.25
	ATOM	81	N	LEU	12	-9.424	50.420	-43.393	1.00	55.34
	ATOM	82	CA	LEU	12	-8.825	51.713	-43.796	1.00	54.13
	ATOM	83	C	LEU	12	-8.546	52.656	-42.612	1.00	54.96
	ATOM	84	O	LEU	12	-7.769	53.603	-42.706	1.00	54.11
20	ATOM	85	CB	LEU	12	-9.707	52.391	-44.864	1.00	54.65
	ATOM	86	CG	LEU	12	-10.141	51.549	-46.077	1.00	20.00
	ATOM	87	CD1	LEU	12	-11.087	52.349	-46.958	1.00	20.00
	ATOM	88	CD2	LEU	12	-8.918	51.093	-46.859	1.00	20.00
	ATOM	89	N	LEU	13	-9.201	52.374	-41.470	1.00	49.64
25	ATOM	90	CA	LEU	13	-9.023	53.164	-40.257	1.00	46.11
	ATOM	91	CB	LEU	13	-10.253	53.195	-39.393	1.00	40.39
	ATOM	92	CG	LEU	13	-9.921	53.965	-38.134	1.00	34.58
	ATOM	93	CD1	LEU	13	-10.204	55.451	-38.332	1.00	33.04
	ATOM	94	CD2	LEU	13	-10.687	53.423	-36.945	1.00	33.82
30	ATOM	95	C	LEU	13	-7.874	52.640	-39.411	1.00	49.82
	ATOM	96	O	LEU	13	-7.548	53.248	-38.395	1.00	50.94
	ATOM	97	N	VAL	14	-7.246	51.529	-39.807	1.00	51.20
	ATOM	98	CA	VAL	14	-6.007	51.177	-39.084	1.00	51.39
	ATOM	99	C	VAL	14	-4.894	51.163	-40.081	1.00	53.87
35	ATOM	100	O	VAL	14	-3.712	51.223	-39.734	1.00	50.27
	ATOM	101	CB	VAL	14	-5.904	49.790	-38.478	1.00	47.82
	ATOM	102	CG1	VAL	14	-4.597	49.650	-37.704	1.00	20.00
	ATOM	103	CG2	VAL	14	-7.098	49.484	-37.581	1.00	20.00
	ATOM	104	N	ARG	15	-5.310	51.042	-41.349	1.00	58.57
40	ATOM	105	CA	ARG	15	-4.465	51.031	-42.519	1.00	64.33
	ATOM	106	C	ARG	15	-3.619	49.754	-42.596	1.00	66.69
	ATOM	107	O	ARG	15	-2.476	49.742	-43.075	1.00	67.11
	ATOM	108	CB	ARG	15	-3.688	52.370	-42.542	1.00	65.68
	ATOM	109	CG	ARG	15	-4.534	53.631	-42.289	1.00	69.20
45	ATOM	110	CD	ARG	15	-4.295	54.732	-43.318	1.00	20.00
	ATOM	111	NE	ARG	15	-4.941	56.026	-42.995	1.00	20.00
	ATOM	112	CZ	ARG	15	-4.834	57.116	-43.748	1.00	20.00
	ATOM	113	NH1	ARG	15	-4.099	57.087	-44.851	1.00	20.00
	ATOM	114	NH2	ARG	15	-5.462	58.229	-43.395	1.00	20.00
50	ATOM	115	N	HIS	16	-4.248	48.677	-42.088	1.00	69.57
	ATOM	116	CA	HIS	16	-3.676	47.342	-42.057	1.00	71.62
	ATOM	117	C	HIS	16	-4.692	46.241	-42.037	1.00	73.99
	ATOM	118	O	HIS	16	-5.853	46.451	-41.713	1.00	75.45
	ATOM	119	CB	HIS	16	-2.915	47.059	-40.787	1.00	69.66
55	ATOM	120	CG	HIS	16	-1.886	48.084	-40.452	1.00	20.00
	ATOM	121	ND1	HIS	16	-0.629	48.034	-41.019	1.00	20.00
	ATOM	122	CD2	HIS	16	-1.907	49.161	-39.627	1.00	20.00
	ATOM	123	CE1	HIS	16	0.077	49.039	-40.557	1.00	20.00
	ATOM	124	NE2	HIS	16	-0.659	49.730	-39.705	1.00	20.00
60	ATOM	125	N	CYS	17	-4.238	45.054	-42.371	1.00	75.75
	ATOM	126	CA	CYS	17	-5.164	43.945	-42.406	1.00	77.01
	ATOM	127	C	CYS	17	-5.629	43.535	-40.996	1.00	75.86
	ATOM	128	O	CYS	17	-4.905	43.646	-40.011	1.00	73.08
	ATOM	129	CB	CYS	17	-4.531	42.764	-43.138	1.00	80.96
65	ATOM	130	SG	CYS	17	-5.462	42.208	-44.606	1.00	86.58
	ATOM	131	N	VAL	18	-6.873	43.058	-40.947	1.00	75.89

	ATOM 132	CA	VAL	18	-7.514	42.612	-39.743	1.00	77.02
	ATOM 133	C	VAL	18	-8.201	41.316	-40.034	1.00	78.19
	ATOM 134	O	VAL	18	-8.575	41.019	-41.158	1.00	77.54
	ATOM 135	CB	VAL	18	-8.560	43.610	-39.175	1.00	76.68
5	ATOM 136	CG1	VAL	18	-9.698	42.871	-38.504	1.00	76.28
	ATOM 137	CG2	VAL	18	-7.902	44.553	-38.190	1.00	76.37
	ATOM 138	N	ALA	19	-8.389	40.545	-38.960	1.00	80.09
	ATOM 139	CA	ALA	19	-9.085	39.216	-38.988	1.00	82.16
	ATOM 140	C	ALA	19	-10.567	39.374	-38.796	1.00	84.28
10	ATOM 141	O	ALA	19	-10.954	40.053	-37.843	1.00	84.12
	ATOM 142	CB	ALA	19	-8.507	38.294	-37.913	1.00	80.00
	ATOM 143	N	CYS	20	-11.440	38.863	-39.619	1.00	87.37
	ATOM 144	CA	CYS	20	-12.814	39.320	-39.328	1.00	88.33
	ATOM 145	CB	CYS	20	-13.740	38.984	-40.503	1.00	86.90
15	ATOM 146	SG	CYS	20	-14.090	40.383	-41.615	1.00	84.56
	ATOM 147	C	CYS	20	-13.364	38.856	-37.978	1.00	89.64
	ATOM 148	O	CYS	20	-14.489	39.189	-37.617	1.00	89.77
	ATOM 149	N	GLY	21	-12.577	38.076	-37.239	1.00	87.37
	ATOM 150	CA	GLY	21	-13.057	37.603	-35.961	1.00	88.33
20	ATOM 151	C	GLY	21	-13.259	38.765	-34.987	1.00	89.64
	ATOM 152	O	GLY	21	-13.872	38.641	-33.930	1.00	89.77
	ATOM 153	N	LEU	22	-12.710	39.931	-35.346	1.00	87.56
	ATOM 154	CA	LEU	22	-12.781	41.133	-34.513	1.00	85.33
	ATOM 155	CB	LEU	22	-11.610	42.058	-34.793	1.00	79.98
25	ATOM 156	CG	LEU	22	-10.610	42.221	-33.659	1.00	77.91
	ATOM 157	CD1	LEU	22	-9.315	42.832	-34.158	1.00	78.13
	ATOM 158	CD2	LEU	22	-11.210	43.066	-32.535	1.00	72.50
	ATOM 159	C	LEU	22	-14.075	41.903	-34.702	1.00	86.05
	ATOM 160	O	LEU	22	-14.708	42.261	-33.717	1.00	88.06
30	ATOM 161	N	LEU	23	-14.468	42.175	-35.972	1.00	85.11
	ATOM 162	CA	LEU	23	-15.660	42.999	-36.204	1.00	83.33
	ATOM 163	C	LEU	23	-16.993	42.256	-36.164	1.00	84.21
	ATOM 164	O	LEU	23	-18.047	42.892	-36.325	1.00	86.28
	ATOM 165	CB	LEU	23	-15.548	43.731	-37.545	1.00	78.93
35	ATOM 166	CG	LEU	23	-14.247	44.395	-37.971	1.00	73.68
	ATOM 167	CD1	LEU	23	-14.267	44.636	-39.465	1.00	20.00
	ATOM 168	CD2	LEU	23	-14.004	45.677	-37.212	1.00	20.00
	ATOM 169	N	ARG	24	-16.995	40.952	-35.933	1.00	86.09
	ATOM 170	CA	ARG	24	-18.301	40.287	-35.953	1.00	85.39
40	ATOM 171	C	ARG	24	-18.935	40.170	-34.586	1.00	84.10
	ATOM 172	O	ARG	24	-20.156	40.068	-34.487	1.00	83.24
	ATOM 173	CB	ARG	24	-18.232	38.946	-36.626	1.00	80.05
	ATOM 174	CG	ARG	24	-16.925	38.218	-36.506	1.00	20.00
	ATOM 175	CD	ARG	24	-17.172	36.762	-36.786	1.00	20.00
45	ATOM 176	NE	ARG	24	-16.682	36.387	-38.087	1.00	20.00
	ATOM 177	CZ	ARG	24	-16.599	35.108	-38.414	1.00	20.00
	ATOM 178	NH1	ARG	24	-16.982	34.174	-37.553	1.00	20.00
	ATOM 179	NH2	ARG	24	-16.135	34.765	-39.598	1.00	20.00
	ATOM 180	N	THR	25	-18.133	40.162	-33.502	1.00	84.61
50	ATOM 181	CA	THR	25	-18.710	40.059	-32.142	1.00	85.07
	ATOM 182	C	THR	25	-19.279	41.426	-31.759	1.00	85.65
	ATOM 183	O	THR	25	-20.247	41.551	-31.018	1.00	86.77
	ATOM 184	CB	THR	25	-17.656	39.515	-31.158	1.00	20.00
	ATOM 185	OG1	THR	25	-17.052	38.354	-31.725	1.00	20.00
55	ATOM 186	CG2	THR	25	-18.303	39.170	-29.828	1.00	20.00
	ATOM 187	N	PRO	26	-18.588	42.466	-32.327	1.00	84.19
	ATOM 188	CA	PRO	26	-19.067	43.910	-32.346	1.00	82.44
	ATOM 189	C	PRO	26	-20.243	44.189	-33.300	1.00	84.32
	ATOM 190	O	PRO	26	-20.882	45.244	-33.289	1.00	81.24
60	ATOM 191	CB	PRO	26	-17.832	44.692	-32.732	1.00	78.00
	ATOM 192	CG	PRO	26	-16.751	43.965	-32.008	1.00	20.00
	ATOM 193	CD	PRO	26	-17.260	42.571	-31.727	1.00	20.00
	ATOM 194	N	ARG	27	-20.496	43.176	-34.165	1.00	87.06
	ATOM 195	CA	ARG	27	-21.440	43.127	-35.315	1.00	88.17
65	ATOM 196	C	ARG	27	-21.210	44.415	-36.157	1.00	88.45
	ATOM 197	O	ARG	27	-22.096	44.901	-36.857	1.00	84.89

	ATOM 198	CB	ARG	27	-22.867	42.645	-34.887	0.00	86.84
	ATOM 199	CG	ARG	27	-22.944	41.084	-34.642	0.00	85.55
	ATOM 200	CD	ARG	27	-24.087	40.235	-35.324	1.00	20.00
	ATOM 201	NE	ARG	27	-24.064	38.772	-35.033	1.00	20.00
5	ATOM 202	CZ	ARG	27	-24.890	37.820	-35.491	1.00	20.00
	ATOM 203	NH1	ARG	27	-25.864	38.154	-36.313	1.00	20.00
	ATOM 204	NH2	ARG	27	-24.735	36.562	-35.111	1.00	20.00
	ATOM 205	N	PRO	28	-19.955	44.891	-36.026	1.00	89.49
	ATOM 206	CA	PRO	28	-19.450	45.984	-36.926	1.00	91.86
10	ATOM 207	C	PRO	28	-19.599	45.567	-38.432	1.00	95.75
	ATOM 208	O	PRO	28	-19.422	46.355	-39.358	1.00	96.59
	ATOM 209	CB	PRO	28	-18.060	46.291	-36.355	0.00	91.23
	ATOM 210	CG	PRO	28	-18.313	46.151	-34.873	1.00	20.00
	ATOM 211	CD	PRO	28	-19.485	45.231	-34.677	1.00	20.00
15	ATOM 212	N	LYS	29	-19.942	44.268	-38.610	1.00	98.41
	ATOM 213	CA	LYS	29	-20.161	43.512	-39.862	1.00	96.39
	ATOM 214	C	LYS	29	-19.582	43.960	-41.199	1.00	96.88
	ATOM 215	O	LYS	29	-20.054	44.925	-41.796	1.00	94.52
	ATOM 216	CB	LYS	29	-21.665	43.308	-40.032	0.00	95.29
20	ATOM 217	CG	LYS	29	-22.511	44.276	-39.233	0.00	92.34
	ATOM 218	CD	LYS	29	-23.066	45.376	-40.114	0.00	89.50
	ATOM 219	CE	LYS	29	-24.335	45.976	-39.521	1.00	20.00
	ATOM 220	NZ	LYS	29	-25.408	46.125	-40.539	1.00	20.00
	ATOM 221	N	PRO	30	-18.490	43.247	-41.652	1.00	97.72
25	ATOM 222	CA	PRO	30	-17.977	43.420	-43.055	1.00	98.94
	ATOM 223	C	PRO	30	-18.914	42.920	-44.160	1.00	100.71
	ATOM 224	O	PRO	30	-19.282	41.738	-44.168	1.00	100.56
	ATOM 225	CB	PRO	30	-16.636	42.734	-43.052	0.00	97.33
	ATOM 226	CG	PRO	30	-16.109	43.061	-41.708	1.00	20.00
30	ATOM 227	CD	PRO	30	-17.288	43.403	-40.828	1.00	20.00
	ATOM 228	N	ALA	31	-19.296	43.791	-45.097	1.00	101.17
	ATOM 229	CA	ALA	31	-20.177	43.392	-46.169	1.00	99.22
	ATOM 230	C	ALA	31	-19.416	43.259	-47.487	1.00	99.01
	ATOM 231	O	ALA	31	-19.894	42.648	-48.446	1.00	102.17
35	ATOM 232	CB	ALA	31	-21.315	44.394	-46.298	1.00	20.00
	ATOM 233	N	GLY	32	-18.219	43.816	-47.478	1.00	95.52
	ATOM 234	CA	GLY	32	-17.356	43.654	-48.594	1.00	91.44
	ATOM 235	C	GLY	32	-16.421	42.510	-48.239	1.00	88.49
	ATOM 236	O	GLY	32	-15.272	42.736	-47.890	1.00	86.70
40	ATOM 237	N	ALA	33	-16.916	41.286	-48.344	1.00	15.00
	ATOM 238	CA	ALA	33	-16.099	40.118	-48.074	1.00	15.00
	ATOM 239	C	ALA	33	-16.316	39.467	-46.730	1.00	15.00
	ATOM 240	O	ALA	33	-16.156	38.253	-46.582	1.00	15.00
	ATOM 241	CB	ALA	33	-14.624	40.489	-48.220	1.00	15.00
45	ATOM 242	N	SER	34	-16.685	40.240	-45.743	1.00	15.00
	ATOM 243	CA	SER	34	-16.803	39.676	-44.397	1.00	15.00
	ATOM 244	C	SER	34	-18.083	38.890	-44.135	1.00	15.00
	ATOM 245	O	SER	34	-18.088	37.908	-43.389	1.00	15.00
	ATOM 246	CB	SER	34	-16.682	40.801	-43.379	1.00	15.00
50	ATOM 247	OG	SER	34	-16.330	40.287	-42.106	1.00	20.00
	ATOM 248	N	SER	35	-19.181	39.308	-44.738	1.00	15.00
	ATOM 249	CA	SER	35	-20.480	38.657	-44.530	1.00	15.00
	ATOM 250	C	SER	35	-20.312	37.138	-44.584	1.00	15.00
	ATOM 251	O	SER	35	-20.909	36.373	-43.827	1.00	15.00
55	ATOM 252	CB	SER	35	-21.508	39.127	-45.572	1.00	15.00
	ATOM 253	OG	SER	35	-21.002	40.220	-46.326	1.00	20.00
	ATOM 254	N	PRO	36	-19.459	36.761	-45.530	1.00	15.00
	ATOM 255	CA	PRO	36	-19.085	35.338	-45.751	1.00	15.00
	ATOM 256	C	PRO	36	-18.688	34.620	-44.490	1.00	15.00
60	ATOM 257	O	PRO	36	-19.237	33.572	-44.150	1.00	15.00
	ATOM 258	CB	PRO	36	-18.106	35.444	-46.920	1.00	15.00
	ATOM 259	CG	PRO	36	-18.714	36.551	-47.744	1.00	15.00
	ATOM 260	CD	PRO	36	-19.506	37.428	-46.839	1.00	15.00
	ATOM 261	N	ALA	37	-17.736	35.208	-43.791	1.00	15.00
65	ATOM 262	CA	ALA	37	-17.275	34.690	-42.521	1.00	15.00
	ATOM 263	C	ALA	37	-18.424	34.697	-41.526	1.00	15.00

	ATOM 264	O	ALA	37	-18.542	33.807	-40.694	1.00	15.00
	ATOM 265	CB	ALA	37	-16.114	35.518	-42.004	1.00	15.00
	ATOM 266	N	PRO	38	-19.264	35.759	-41.594	1.00	15.00
	ATOM 267	CA	PRO	38	-20.428	35.878	-40.657	1.00	15.00
5	ATOM 268	C	PRO	38	-21.308	34.692	-40.558	1.00	15.00
	ATOM 269	O	PRO	38	-21.590	34.168	-39.490	1.00	15.00
	ATOM 270	CB	PRO	38	-21.130	37.129	-41.082	1.00	15.00
	ATOM 271	CG	PRO	38	-19.951	38.007	-41.388	1.00	20.00
	ATOM 272	CD	PRO	38	-18.778	37.127	-41.761	1.00	20.00
10	ATOM 273	N	ARG	39	-21.734	34.261	-41.737	1.00	15.00
	ATOM 274	CA	ARG	39	-22.628	33.108	-41.839	1.00	15.00
	ATOM 275	C	ARG	39	-21.850	31.777	-41.757	1.00	15.00
	ATOM 276	O	ARG	39	-22.468	30.711	-41.829	1.00	15.00
	ATOM 277	CB	ARG	39	-23.434	33.145	-43.119	1.00	15.00
15	ATOM 278	CG	ARG	39	-24.164	34.437	-43.355	1.00	15.00
	ATOM 279	CD	ARG	39	-24.079	34.778	-44.825	1.00	15.00
	ATOM 280	NE	ARG	39	-24.795	35.996	-45.141	1.00	20.00
	ATOM 281	CZ	ARG	39	-25.402	36.210	-46.303	1.00	20.00
	ATOM 282	NH1	ARG	39	-25.393	35.269	-47.238	1.00	20.00
20	ATOM 283	NH2	ARG	39	-26.018	37.361	-46.528	1.00	20.00
	ATOM 284	N	THR	40	-20.521	31.849	-41.646	1.00	15.00
	ATOM 285	CA	THR	40	-19.589	30.692	-41.579	1.00	15.00
	ATOM 286	C	THR	40	-19.458	30.229	-40.125	1.00	15.00
	ATOM 287	O	THR	40	-19.577	29.061	-39.769	1.00	15.00
25	ATOM 288	CB	THR	40	-18.213	31.065	-42.079	1.00	15.00
	ATOM 289	OG1	THR	40	-18.363	31.800	-43.305	1.00	20.00
	ATOM 290	CG2	THR	40	-17.378	29.824	-42.323	1.00	20.00
	ATOM 291	N	ALA	41	-19.210	31.240	-39.297	1.00	15.00
	ATOM 292	CA	ALA	41	-19.125	31.016	-37.881	1.00	15.00
30	ATOM 293	C	ALA	41	-20.522	30.931	-37.274	1.00	15.00
	ATOM 294	O	ALA	41	-20.671	30.462	-36.160	1.00	15.00
	ATOM 295	CB	ALA	41	-18.335	32.122	-37.190	1.00	15.00
	ATOM 296	N	LEU	42	-21.523	31.385	-37.976	1.00	15.00
	ATOM 297	CA	LEU	42	-22.896	31.324	-37.490	1.00	15.00
35	ATOM 298	C	LEU	42	-23.440	29.918	-37.696	1.00	15.00
	ATOM 299	O	LEU	42	-24.048	29.330	-36.796	1.00	15.00
	ATOM 300	CB	LEU	42	-23.780	32.322	-38.219	1.00	15.00
	ATOM 301	CG	LEU	42	-23.514	33.804	-37.952	1.00	20.00
	ATOM 302	CD1	LEU	42	-24.496	34.675	-38.729	1.00	20.00
40	ATOM 303	CD2	LEU	42	-23.588	34.104	-36.469	1.00	20.00
	ATOM 304	N	GLN	43	-23.214	29.367	-38.878	1.00	15.00
	ATOM 305	CA	GLN	43	-23.619	27.991	-39.212	1.00	15.00
	ATOM 306	C	GLN	43	-22.992	27.013	-38.247	1.00	15.00
	ATOM 307	O	GLN	43	-23.682	26.319	-37.504	1.00	15.00
45	ATOM 308	CB	GLN	43	-23.200	27.617	-40.648	1.00	15.00
	ATOM 309	CG	GLN	43	-23.510	28.629	-41.725	1.00	20.00
	ATOM 310	CD	GLN	43	-22.948	28.156	-43.050	1.00	20.00
	ATOM 311	OE1	GLN	43	-22.253	27.147	-43.132	1.00	20.00
	ATOM 312	NE2	GLN	43	-23.151	28.742	-44.227	1.00	20.00
50	END				31.712	6.654	-112.989	0.00	0.00

TABLE 24

	ATOM	2	CA	PRO	1	11.710	58.647	-8.623	1.00	98.53
	ATOM	3	C	PRO	1	10.938	59.150	-7.465	1.00	101.54
5	ATOM	4	O	PRO	1	11.579	59.741	-6.585	1.00	102.94
	ATOM	5	CB	PRO	1	13.194	58.569	-8.320	1.00	96.91
	ATOM	6	CG	PRO	1	13.739	58.461	-9.698	1.00	20.00
	ATOM	7	CD	PRO	1	12.708	58.994	-10.680	1.00	20.00
	ATOM	8	N	THR	2	9.666	59.031	-7.330	1.00	104.82
10	ATOM	9	CA	THR	2	9.310	59.574	-6.060	1.00	107.58
	ATOM	10	C	THR	2	9.002	58.400	-5.145	1.00	108.57
	ATOM	11	O	THR	2	9.559	57.327	-5.378	1.00	109.82
	ATOM	12	CB	THR	2	8.219	60.622	-6.191	1.00	20.00
	ATOM	13	OG1	THR	2	8.820	61.911	-6.450	1.00	20.00
15	ATOM	14	CG2	THR	2	7.386	60.672	-4.920	1.00	20.00
	ATOM	15	N	PRO	3	8.155	58.502	-4.094	1.00	108.52
	ATOM	16	CA	PRO	3	8.105	57.328	-3.181	1.00	109.56
	ATOM	17	C	PRO	3	7.954	55.965	-3.767	1.00	110.79
	ATOM	18	O	PRO	3	7.008	55.736	-4.520	1.00	114.50
20	ATOM	19	CB	PRO	3	7.031	57.699	-2.179	1.00	108.23
	ATOM	20	CG	PRO	3	7.318	59.150	-1.997	1.00	109.80
	ATOM	21	CD	PRO	3	8.021	59.671	-3.235	1.00	110.35
	ATOM	22	C	CYS	4	8.610	52.487	-3.575	1.00	101.54
	ATOM	23	O	CYS	4	9.744	52.005	-3.463	1.00	102.94
25	ATOM	24	CB	CYS	4	8.728	53.940	-5.612	1.00	96.91
	ATOM	25	SG	CYS	4	7.316	54.077	-6.754	1.00	92.39
	ATOM	26	N	CYS	4	8.796	55.004	-3.457	1.00	93.44
	ATOM	27	CA	CYS	4	8.317	53.837	-4.187	1.00	98.53
	ATOM	28	N	VAL	5	7.483	51.910	-3.232	1.00	104.82
30	ATOM	29	CA	VAL	5	7.379	50.630	-2.577	1.00	107.58
	ATOM	30	C	VAL	5	7.989	49.468	-3.410	1.00	108.57
	ATOM	31	O	VAL	5	7.839	49.438	-4.635	1.00	109.82
	ATOM	32	CB	VAL	5	5.910	50.521	-2.139	1.00	109.37
	ATOM	33	CG1	VAL	5	5.737	49.389	-1.137	1.00	20.00
35	ATOM	34	CG2	VAL	5	5.405	51.833	-1.567	1.00	20.00
	ATOM	35	N	PRO	6	8.684	48.535	-2.724	1.00	108.52
	ATOM	36	CA	PRO	6	9.313	47.453	-3.515	1.00	109.56
	ATOM	37	C	PRO	6	8.403	46.783	-4.555	1.00	110.79
	ATOM	38	O	PRO	6	7.247	46.506	-4.268	1.00	114.50
40	ATOM	39	CB	PRO	6	9.968	46.586	-2.451	1.00	108.23
	ATOM	40	CG	PRO	6	10.473	47.581	-1.473	1.00	109.80
	ATOM	41	CD	PRO	6	9.625	48.826	-1.617	1.00	110.35
	ATOM	42	N	ALA	7	8.915	46.502	-5.751	1.00	109.87
	ATOM	43	CA	ALA	7	8.128	45.939	-6.847	1.00	107.78
45	ATOM	44	C	ALA	7	7.604	47.014	-7.820	1.00	105.15
	ATOM	45	O	ALA	7	7.688	46.805	-9.018	1.00	103.42
	ATOM	46	CB	ALA	7	6.973	45.123	-6.304	1.00	109.08
	ATOM	47	N	GLU	8	7.100	48.204	-7.360	1.00	101.11
	ATOM	48	CA	GLU	8	6.496	49.197	-8.338	1.00	95.31
50	ATOM	49	CB	GLU	8	5.240	49.838	-7.807	1.00	92.47
	ATOM	50	CG	GLU	8	4.719	49.232	-6.517	1.00	86.82
	ATOM	51	CD	GLU	8	4.355	50.298	-5.497	1.00	85.55
	ATOM	52	OE1	GLU	8	5.220	51.148	-5.182	1.00	85.72
	ATOM	53	OE2	GLU	8	3.207	50.297	-5.031	1.00	81.01
55	ATOM	54	C	GLU	8	7.365	50.327	-8.854	1.00	92.66
	ATOM	55	O	GLU	8	8.421	50.641	-8.292	1.00	94.43
	ATOM	56	N	CYS	9	6.900	50.943	-9.935	1.00	85.19
	ATOM	57	CA	CYS	9	7.635	52.020	-10.580	1.00	78.14
	ATOM	58	C	CYS	9	6.719	53.218	-10.721	1.00	75.43
60	ATOM	59	O	CYS	9	5.551	53.083	-11.078	1.00	76.93
	ATOM	60	CB	CYS	9	8.143	51.586	-11.954	1.00	76.59
	ATOM	61	SG	CYS	9	6.839	51.326	-13.199	1.00	20.00
	ATOM	62	N	PHE	10	7.286	54.387	-10.427	1.00	69.11
	ATOM	63	CA	PHE	10	6.592	55.693	-10.541	1.00	62.54
65	ATOM	64	CB	PHE	10	7.498	56.849	-9.965	1.00	61.14
	ATOM	65	CG	PHE	10	6.854	58.217	-9.958	1.00	61.11

	ATOM	66	CD1	PHE	10	5.819	58.472	-9.065	1.00	59.90
	ATOM	67	CD2	PHE	10	7.234	59.218	-10.852	1.00	63.23
	ATOM	68	CE1	PHE	10	5.170	59.707	-9.063	1.00	61.45
	ATOM	69	CE2	PHE	10	6.594	60.463	-10.864	1.00	61.15
5	ATOM	70	CZ	PHE	10	5.561	60.709	-9.971	1.00	61.89
	ATOM	71	C	PHE	10	6.240	55.970	-12.013	1.00	60.79
	ATOM	72	O	PHE	10	7.125	56.075	-12.869	1.00	63.10
	ATOM	73	N	ASP	11	4.942	56.069	-12.326	1.00	57.06
	ATOM	74	CA	ASP	11	4.513	56.376	-13.707	1.00	52.20
10	ATOM	75	CB	ASP	11	3.138	55.685	-14.044	1.00	46.52
	ATOM	76	CG	ASP	11	2.635	55.837	-15.490	1.00	47.00
	ATOM	77	OD1	ASP	11	2.924	56.891	-16.099	1.00	49.00
	ATOM	78	OD2	ASP	11	1.973	54.904	-15.990	1.00	48.43
	ATOM	79	C	ASP	11	4.372	57.908	-13.816	1.00	53.02
15	ATOM	80	O	ASP	11	3.556	58.481	-13.102	1.00	53.25
	ATOM	81	N	LEU	12	5.145	58.579	-14.690	1.00	55.34
	ATOM	82	CA	LEU	12	5.074	60.049	-14.861	1.00	54.13
	ATOM	83	C	LEU	12	3.808	60.520	-15.598	1.00	54.96
	ATOM	84	O	LEU	12	3.419	61.684	-15.541	1.00	54.11
20	ATOM	85	CB	LEU	12	6.355	60.560	-15.552	1.00	54.65
	ATOM	86	CG	LEU	12	7.709	60.107	-14.979	1.00	20.00
	ATOM	87	CD1	LEU	12	8.844	60.619	-15.850	1.00	20.00
	ATOM	88	CD2	LEU	12	7.858	60.593	-13.545	1.00	20.00
	ATOM	89	N	LEU	13	3.153	59.578	-16.303	1.00	49.64
25	ATOM	90	CA	LEU	13	1.927	59.868	-17.036	1.00	46.11
	ATOM	91	CB	LEU	13	1.756	59.007	-18.256	1.00	40.39
	ATOM	92	CG	LEU	13	0.417	59.337	-18.878	1.00	34.58
	ATOM	93	CD1	LEU	13	0.567	60.474	-19.885	1.00	33.04
	ATOM	94	CD2	LEU	13	-0.198	58.116	-19.530	1.00	33.82
30	ATOM	95	C	LEU	13	0.696	59.686	-16.162	1.00	49.82
	ATOM	96	O	LEU	13	-0.411	59.970	-16.614	1.00	50.94
	ATOM	97	N	VAL	14	0.859	59.226	-14.919	1.00	51.20
	ATOM	98	CA	VAL	14	-0.325	59.269	-14.037	1.00	51.39
	ATOM	99	C	VAL	14	0.016	60.137	-12.869	1.00	53.87
35	ATOM	100	O	VAL	14	-0.856	60.619	-12.143	1.00	50.27
	ATOM	101	CB	VAL	14	-0.756	57.982	-13.358	1.00	47.82
	ATOM	102	CG1	VAL	14	-2.040	58.211	-12.566	1.00	20.00
	ATOM	103	CG2	VAL	14	-0.936	56.854	-14.367	1.00	20.00
	ATOM	104	N	ARG	15	1.334	60.293	-12.682	1.00	58.57
40	ATOM	105	CA	ARG	15	1.954	61.099	-11.657	1.00	64.33
	ATOM	106	C	ARG	15	1.753	60.494	-10.263	1.00	66.69
	ATOM	107	O	ARG	15	1.627	61.193	-9.246	1.00	67.11
	ATOM	108	CB	ARG	15	1.462	62.554	-11.851	1.00	65.68
	ATOM	109	CG	ARG	15	1.510	63.084	-13.295	1.00	69.20
45	ATOM	110	CD	ARG	15	2.178	64.451	-13.413	1.00	20.00
	ATOM	111	NE	ARG	15	2.067	65.076	-14.752	1.00	20.00
	ATOM	112	CZ	ARG	15	2.557	66.273	-15.053	1.00	20.00
	ATOM	113	NH1	ARG	15	3.173	66.989	-14.121	1.00	20.00
	ATOM	114	NH2	ARG	15	2.431	66.751	-16.283	1.00	20.00
50	ATOM	115	N	HIS	16	1.723	59.148	-10.274	1.00	69.57
	ATOM	116	CA	HIS	16	1.565	58.328	-9.085	1.00	71.62
	ATOM	117	C	HIS	16	2.149	56.954	-9.205	1.00	73.99
	ATOM	118	O	HIS	16	2.396	56.462	-10.298	1.00	75.45
	ATOM	119	CB	HIS	16	0.124	58.007	-8.781	1.00	69.66
55	ATOM	120	CG	HIS	16	-0.768	59.201	-8.724	1.00	20.00
	ATOM	121	ND1	HIS	16	-0.867	59.954	-7.573	1.00	20.00
	ATOM	122	CD2	HIS	16	-1.592	59.767	-9.642	1.00	20.00
	ATOM	123	CE1	HIS	16	-1.713	60.934	-7.787	1.00	20.00
	ATOM	124	NE2	HIS	16	-2.178	60.842	-9.020	1.00	20.00
60	ATOM	125	N	CYS	17	2.349	56.328	-8.068	1.00	75.75
	ATOM	126	CA	CYS	17	2.938	55.009	-8.099	1.00	77.01
	ATOM	127	C	CYS	17	1.974	53.958	-8.679	1.00	75.86
	ATOM	128	O	CYS	17	0.757	54.039	-8.536	1.00	73.08
	ATOM	129	CB	CYS	17	3.401	54.615	-6.698	1.00	80.96
65	ATOM	130	SG	CYS	17	5.185	54.249	-6.575	1.00	86.58
	ATOM	131	N	VAL	18	2.574	52.970	-9.341	1.00	75.89

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	ATOM	132	CA	VAL	18	1.876	51.880	-9.961	1.00	77.02
	ATOM	133	C	VAL	18	2.595	50.613	-9.620	1.00	78.19
	ATOM	134	O	VAL	18	3.786	50.597	-9.349	1.00	77.54
	ATOM	135	CB	VAL	18	1.774	51.990	-11.506	1.00	76.68
5	ATOM	136	CG1	VAL	18	1.808	50.619	-12.147	1.00	76.28
	ATOM	137	CG2	VAL	18	0.500	52.711	-11.896	1.00	76.37
	ATOM	138	N	ALA	19	1.829	49.520	-9.672	1.00	80.09
	ATOM	139	CA	ALA	19	2.326	48.130	-9.407	1.00	82.16
	ATOM	140	C	ALA	19	2.848	47.490	-10.663	1.00	84.28
10	ATOM	141	O	ALA	19	2.129	47.517	-11.664	1.00	84.12
	ATOM	142	CB	ALA	19	1.210	47.275	-8.803	1.00	80.00
	ATOM	143	N	CYS	20	4.037	46.959	-10.739	1.00	87.37
	ATOM	144	CA	CYS	20	4.389	46.577	-12.121	1.00	88.33
	ATOM	145	CB	CYS	20	5.892	46.287	-12.219	1.00	86.90
15	ATOM	146	SG	CYS	20	6.880	47.646	-12.919	1.00	84.56
	ATOM	147	C	CYS	20	3.522	45.464	-12.712	1.00	89.64
	ATOM	148	O	CYS	20	3.709	45.074	-13.861	1.00	89.77
	ATOM	149	N	GLY	21	2.585	44.942	-11.922	1.00	87.37
	ATOM	150	CA	GLY	21	1.749	43.880	-12.432	1.00	88.33
20	ATOM	151	C	GLY	21	0.872	44.377	-13.582	1.00	89.64
	ATOM	152	O	GLY	21	0.255	43.612	-14.320	1.00	89.77
	ATOM	153	N	LEU	22	0.800	45.704	-13.732	1.00	87.56
	ATOM	154	CA	LEU	22	-0.020	46.345	-14.761	1.00	85.33
	ATOM	155	CB	LEU	22	-0.432	47.742	-14.336	1.00	79.98
25	ATOM	156	CG	LEU	22	-1.916	47.942	-14.069	1.00	77.91
	ATOM	157	CD1	LEU	22	-2.162	49.223	-13.296	1.00	78.13
	ATOM	158	CD2	LEU	22	-2.700	47.942	-15.381	1.00	72.50
	ATOM	159	C	LEU	22	0.680	46.424	-16.106	1.00	86.05
	ATOM	160	O	LEU	22	0.085	46.064	-17.113	1.00	88.06
30	ATOM	161	N	LEU	23	1.947	46.909	-16.130	1.00	85.11
	ATOM	162	CA	LEU	23	2.630	47.095	-17.415	1.00	83.33
	ATOM	163	C	LEU	23	3.309	45.853	-17.986	1.00	84.21
	ATOM	164	O	LEU	23	3.885	45.926	-19.084	1.00	86.28
	ATOM	165	CB	LEU	23	3.669	48.216	-17.308	1.00	78.93
35	ATOM	166	CG	LEU	23	3.351	49.516	-16.586	1.00	73.68
	ATOM	167	CD1	LEU	23	4.638	50.232	-16.240	1.00	20.00
	ATOM	168	CD2	LEU	23	2.437	50.395	-17.404	1.00	20.00
	ATOM	169	N	ARG	24	3.247	44.717	-17.308	1.00	86.09
	ATOM	170	CA	ARG	24	3.957	43.572	-17.885	1.00	85.39
40	ATOM	171	C	ARG	24	3.080	42.693	-18.744	1.00	84.10
	ATOM	172	O	ARG	24	3.586	42.000	-19.625	1.00	83.24
	ATOM	173	CB	ARG	24	4.654	42.762	-16.829	1.00	80.05
	ATOM	174	CG	ARG	24	4.004	42.747	-15.477	1.00	20.00
	ATOM	175	CD	ARG	24	4.520	41.555	-14.720	1.00	20.00
45	ATOM	176	NE	ARG	24	5.461	41.945	-13.704	1.00	20.00
	ATOM	177	CZ	ARG	24	5.842	41.069	-12.790	1.00	20.00
	ATOM	178	NH1	ARG	24	5.373	39.828	-12.820	1.00	20.00
	ATOM	179	NH2	ARG	24	6.691	41.431	-11.850	1.00	20.00
	ATOM	180	N	THR	25	1.752	42.679	-18.508	1.00	84.61
50	ATOM	181	CA	THR	25	0.851	41.841	-19.332	1.00	85.07
	ATOM	182	C	THR	25	0.644	42.539	-20.676	1.00	85.65
	ATOM	183	O	THR	25	0.445	41.921	-21.715	1.00	86.77
	ATOM	184	CB	THR	25	-0.452	41.549	-18.563	1.00	20.00
	ATOM	185	OG1	THR	25	-0.123	41.098	-17.251	1.00	20.00
55	ATOM	186	CG2	THR	25	-1.268	40.492	-19.289	1.00	20.00
	ATOM	187	N	PRO	26	0.700	43.907	-20.577	1.00	84.19
	ATOM	188	CA	PRO	26	0.792	44.854	-21.765	1.00	82.44
	ATOM	189	C	PRO	26	2.155	44.864	-22.480	1.00	84.32
	ATOM	190	O	PRO	26	2.339	45.410	-23.570	1.00	81.24
60	ATOM	191	CB	PRO	26	0.457	46.204	-21.173	1.00	78.00
	ATOM	192	CG	PRO	26	-0.613	45.869	-20.190	1.00	20.00
	ATOM	193	CD	PRO	26	-0.468	44.404	-19.854	1.00	20.00
	ATOM	194	N	ARG	27	3.138	44.234	-21.790	1.00	87.06
	ATOM	195	CA	ARG	27	4.597	44.159	-22.076	1.00	88.17
65	ATOM	196	C	ARG	27	5.085	45.606	-22.372	1.00	88.45
	ATOM	197	O	ARG	27	6.068	45.829	-23.077	1.00	84.89

	ATOM	198	CB	ARG	27	4.955	42.945	-22.999	0.00	86.84
	ATOM	199	CG	ARG	27	4.943	41.561	-22.232	0.00	85.55
	ATOM	200	CD	ARG	27	6.173	40.580	-22.358	1.00	20.00
	ATOM	201	NE	ARG	27	6.062	39.307	-21.589	1.00	20.00
5	ATOM	202	CZ	ARG	27	6.957	38.312	-21.496	1.00	20.00
	ATOM	203	NH1	ARG	27	8.103	38.414	-22.137	1.00	20.00
	ATOM	204	NH2	ARG	27	6.684	37.234	-20.777	1.00	20.00
	ATOM	205	N	PRO	28	4.323	46.535	-21.761	1.00	89.49
	ATOM	206	CA	PRO	28	4.751	47.976	-21.719	1.00	91.86
10	ATOM	207	C	PRO	28	6.180	48.103	-21.084	1.00	95.75
	ATOM	208	O	PRO	28	6.820	49.153	-21.099	1.00	96.59
	ATOM	209	CB	PRO	28	3.558	48.675	-21.058	0.00	91.23
	ATOM	210	CG	PRO	28	2.400	47.916	-21.660	1.00	20.00
	ATOM	211	CD	PRO	28	2.885	46.551	-22.061	1.00	20.00
15	ATOM	212	N	LYS	29	6.637	46.957	-20.524	1.00	98.41
	ATOM	213	CA	LYS	29	7.914	46.688	-19.828	1.00	96.39
	ATOM	214	C	LYS	29	8.757	47.797	-19.211	1.00	96.88
	ATOM	215	O	LYS	29	9.401	48.566	-19.922	1.00	94.52
	ATOM	216	CB	LYS	29	8.801	45.875	-20.769	0.00	95.29
20	ATOM	217	CG	LYS	29	8.404	45.974	-22.227	0.00	92.34
	ATOM	218	CD	LYS	29	9.322	46.913	-22.982	0.00	89.50
	ATOM	219	CE	LYS	29	9.345	46.589	-24.470	1.00	20.00
	ATOM	220	NZ	LYS	29	10.729	46.565	-25.012	1.00	20.00
	ATOM	221	N	PRO	30	8.708	47.898	-17.835	1.00	97.72
25	ATOM	222	CA	PRO	30	9.669	48.778	-17.084	1.00	98.94
	ATOM	223	C	PRO	30	11.133	48.326	-17.119	1.00	100.71
	ATOM	224	O	PRO	30	11.440	47.201	-16.704	1.00	100.56
	ATOM	225	CB	PRO	30	9.100	48.855	-15.692	0.00	97.33
	ATOM	226	CG	PRO	30	7.641	48.889	-15.941	1.00	20.00
30	ATOM	227	CD	PRO	30	7.399	48.298	-17.310	1.00	20.00
	ATOM	228	N	ALA	31	12.040	49.181	-17.596	1.00	101.17
	ATOM	229	CA	ALA	31	13.438	48.826	-17.658	1.00	99.22
	ATOM	230	C	ALA	31	14.240	49.545	-16.574	1.00	99.01
	ATOM	231	O	ALA	31	15.369	49.168	-16.254	1.00	102.17
35	ATOM	232	CB	ALA	31	13.987	49.144	-19.041	1.00	20.00
	ATOM	233	N	GLY	32	13.603	50.556	-16.013	1.00	95.52
	ATOM	234	CA	GLY	32	14.182	51.229	-14.905	1.00	91.44
	ATOM	235	C	GLY	32	13.547	50.620	-13.666	1.00	88.49
	ATOM	236	O	GLY	32	12.671	51.219	-13.061	1.00	86.70
40	ATOM	237	N	ALA	33	14.005	49.436	-13.287	1.00	15.00
	ATOM	238	CA	ALA	33	13.503	48.782	-12.093	1.00	15.00
	ATOM	239	C	ALA	33	12.502	47.676	-12.326	1.00	15.00
	ATOM	240	O	ALA	33	12.426	46.719	-11.552	1.00	15.00
	ATOM	241	CB	ALA	33	12.887	49.828	-11.165	1.00	15.00
45	ATOM	242	N	SER	34	11.736	47.775	-13.379	1.00	15.00
	ATOM	243	CA	SER	34	10.677	46.786	-13.590	1.00	15.00
	ATOM	244	C	SER	34	11.142	45.455	-14.171	1.00	15.00
	ATOM	245	O	SER	34	10.597	44.395	-13.854	1.00	15.00
	ATOM	246	CB	SER	34	9.612	47.389	-14.494	1.00	15.00
50	ATOM	247	OG	SER	34	8.388	46.687	-14.361	1.00	20.00
	ATOM	248	N	SER	35	12.147	45.490	-15.027	1.00	15.00
	ATOM	249	CA	SER	35	12.654	44.278	-15.683	1.00	15.00
	ATOM	250	C	SER	35	12.782	43.150	-14.658	1.00	15.00
	ATOM	251	O	SER	35	12.487	41.982	-14.910	1.00	15.00
55	ATOM	252	CB	SER	35	14.004	44.544	-16.369	1.00	15.00
	ATOM	253	OG	SER	35	14.304	45.932	-16.376	1.00	20.00
	ATOM	254	N	PRO	36	13.241	43.584	-13.489	1.00	15.00
	ATOM	255	CA	PRO	36	13.406	42.690	-12.311	1.00	15.00
	ATOM	256	C	PRO	36	12.192	41.849	-12.022	1.00	15.00
60	ATOM	257	O	PRO	36	12.268	40.623	-11.935	1.00	15.00
	ATOM	258	CB	PRO	36	13.948	43.653	-11.254	1.00	15.00
	ATOM	259	CG	PRO	36	14.840	44.553	-12.073	1.00	15.00
	ATOM	260	CD	PRO	36	14.334	44.566	-13.473	1.00	15.00
	ATOM	261	N	ALA	37	11.066	42.525	-11.891	1.00	15.00
65	ATOM	262	CA	ALA	37	9.793	41.872	-11.674	1.00	15.00
	ATOM	263	C	ALA	37	9.472	40.983	-12.864	1.00	15.00

	ATOM	264	O	ALA	37	8.896	39.913	-12.712	1.00	15.00
	ATOM	265	CB	ALA	37	8.701	42.904	-11.466	1.00	15.00
	ATOM	266	N	PRO	38	9.819	41.468	-14.080	1.00	15.00
	ATOM	267	CA	PRO	38	9.544	40.683	-15.327	1.00	15.00
5	ATOM	268	C	PRO	38	10.003	39.276	-15.325	1.00	15.00
	ATOM	269	O	PRO	38	9.260	38.340	-15.587	1.00	15.00
	ATOM	270	CB	PRO	38	10.117	41.513	-16.432	1.00	15.00
	ATOM	271	CG	PRO	38	9.729	42.886	-15.964	1.00	20.00
	ATOM	272	CD	PRO	38	9.588	42.861	-14.458	1.00	20.00
10	ATOM	273	N	ARG	39	11.281	39.146	-14.995	1.00	15.00
	ATOM	274	CA	ARG	39	11.918	37.831	-14.955	1.00	15.00
	ATOM	275	C	ARG	39	11.615	37.093	-13.632	1.00	15.00
	ATOM	276	O	ARG	39	12.086	35.967	-13.448	1.00	15.00
	ATOM	277	CB	ARG	39	13.414	37.935	-15.146	1.00	15.00
15	ATOM	278	CG	ARG	39	13.832	38.718	-16.359	1.00	15.00
	ATOM	279	CD	ARG	39	15.038	39.556	-16.002	1.00	15.00
	ATOM	280	NE	ARG	39	15.526	40.313	-17.134	1.00	20.00
	ATOM	281	CZ	ARG	39	16.807	40.613	-17.320	1.00	20.00
	ATOM	282	NH1	ARG	39	17.718	40.190	-16.454	1.00	20.00
20	ATOM	283	NH2	ARG	39	17.175	41.332	-18.370	1.00	20.00
	ATOM	284	N	THR	40	10.878	37.738	-12.725	1.00	15.00
	ATOM	285	CA	THR	40	10.498	37.220	-11.385	1.00	15.00
	ATOM	286	C	THR	40	9.215	36.390	-11.505	1.00	15.00
	ATOM	287	O	THR	40	9.085	35.266	-11.034	1.00	15.00
25	ATOM	288	CB	THR	40	10.238	38.347	-10.413	1.00	15.00
	ATOM	289	OG1	THR	40	11.302	39.305	-10.540	1.00	20.00
	ATOM	290	CG2	THR	40	10.185	37.826	-8.990	1.00	20.00
	ATOM	291	N	ALA	41	8.268	37.030	-12.186	1.00	15.00
	ATOM	292	CA	ALA	41	7.015	36.386	-12.466	1.00	15.00
30	ATOM	293	C	ALA	41	7.161	35.442	-13.657	1.00	15.00
	ATOM	294	O	ALA	41	6.309	34.597	-13.869	1.00	15.00
	ATOM	295	CB	ALA	41	5.918	37.405	-12.754	1.00	15.00
	ATOM	296	N	LEU	42	8.202	35.587	-14.429	1.00	15.00
	ATOM	297	CA	LEU	42	8.439	34.718	-15.575	1.00	15.00
35	ATOM	298	C	LEU	42	9.027	33.400	-15.093	1.00	15.00
	ATOM	299	O	LEU	42	8.594	32.318	-15.504	1.00	15.00
	ATOM	300	CB	LEU	42	9.391	35.366	-16.566	1.00	15.00
	ATOM	301	CG	LEU	42	8.874	36.593	-17.318	1.00	20.00
	ATOM	302	CD1	LEU	42	9.928	37.109	-18.293	1.00	20.00
40	ATOM	303	CD2	LEU	42	7.584	36.273	-18.044	1.00	20.00
	ATOM	304	N	GLN	43	10.009	33.481	-14.209	1.00	15.00
	ATOM	305	CA	GLN	43	10.639	32.299	-13.599	1.00	15.00
	ATOM	306	C	GLN	43	9.602	31.462	-12.889	1.00	15.00
	ATOM	307	O	GLN	43	9.356	30.312	-13.246	1.00	15.00
45	ATOM	308	CB	GLN	43	11.732	32.705	-12.589	1.00	15.00
	ATOM	309	CG	GLN	43	12.712	33.758	-13.048	1.00	20.00
	ATOM	310	CD	GLN	43	13.651	34.113	-11.912	1.00	20.00
	ATOM	311	OE1	GLN	43	13.498	33.654	-10.783	1.00	20.00
	ATOM	312	NE2	GLN	43	14.713	34.908	-12.008	1.00	20.00
50	END					50.903	67.374	64.558	0.00	0.00

TABLE 25

	ATOM	2	CA	PRO	1	-36.025	62.959	-15.214	1.00	98.53
	ATOM	3	C	PRO	1	-36.724	62.591	-16.465	1.00	101.54
5	ATOM	4	O	PRO	1	-37.860	63.060	-16.624	1.00	102.94
	ATOM	5	CB	PRO	1	-36.984	63.490	-14.167	1.00	96.91
	ATOM	6	CG	PRO	1	-36.026	64.147	-13.241	1.00	20.00
	ATOM	7	CD	PRO	1	-34.736	64.438	-13.990	1.00	20.00
	ATOM	8	N	THR	2	-36.227	61.846	-17.386	1.00	104.82
10	ATOM	9	CA	THR	2	-37.227	61.667	-18.388	1.00	107.58
	ATOM	10	C	THR	2	-37.756	60.250	-18.240	1.00	108.57
	ATOM	11	O	THR	2	-37.703	59.729	-17.125	1.00	109.82
	ATOM	12	CB	THR	2	-36.706	62.044	-19.764	1.00	20.00
	ATOM	13	OG1	THR	2	-36.901	63.460	-19.980	1.00	20.00
15	ATOM	14	CG2	THR	2	-37.428	61.241	-20.834	1.00	20.00
	ATOM	15	N	PRO	3	-38.285	59.561	-19.278	1.00	108.52
	ATOM	16	CA	PRO	3	-38.935	58.267	-18.937	1.00	109.56
	ATOM	17	C	PRO	3	-38.207	57.301	-18.063	1.00	110.79
	ATOM	18	O	PRO	3	-37.076	56.937	-18.382	1.00	114.50
20	ATOM	19	CB	PRO	3	-39.342	57.704	-20.283	1.00	108.23
	ATOM	20	CG	PRO	3	-39.790	58.947	-20.973	1.00	109.80
	ATOM	21	CD	PRO	3	-39.096	60.137	-20.342	1.00	110.35
	ATOM	22	C	CYS	4	-38.317	54.736	-15.619	1.00	101.54
	ATOM	23	O	CYS	4	-38.902	54.844	-14.535	1.00	102.94
25	ATOM	24	CB	CYS	4	-36.747	56.686	-15.699	1.00	96.91
	ATOM	25	SG	CYS	4	-35.094	56.533	-16.447	1.00	92.39
	ATOM	26	N	CYS	4	-38.775	56.814	-16.982	1.00	93.44
	ATOM	27	CA	CYS	4	-37.786	55.904	-16.418	1.00	98.53
	ATOM	28	N	VAL	5	-38.023	53.617	-16.235	1.00	104.82
30	ATOM	29	CA	VAL	5	-38.411	52.303	-15.788	1.00	107.58
	ATOM	30	C	VAL	5	-37.848	51.947	-14.383	1.00	108.57
	ATOM	31	O	VAL	5	-36.703	52.286	-14.069	1.00	109.82
	ATOM	32	CB	VAL	5	-38.086	51.366	-16.962	1.00	109.37
	ATOM	33	CG1	VAL	5	-38.760	50.016	-16.766	1.00	20.00
35	ATOM	34	CG2	VAL	5	-38.486	51.984	-18.290	1.00	20.00
	ATOM	35	N	PRO	6	-38.678	51.280	-13.553	1.00	108.52
	ATOM	36	CA	PRO	6	-38.170	50.984	-12.194	1.00	109.56
	ATOM	37	C	PRO	6	-36.759	50.381	-12.136	1.00	110.79
	ATOM	38	O	PRO	6	-36.433	49.509	-12.929	1.00	114.50
40	ATOM	39	CB	PRO	6	-39.320	50.217	-11.557	1.00	108.23
	ATOM	40	CG	PRO	6	-40.519	50.913	-12.085	1.00	109.80
	ATOM	41	CD	PRO	6	-40.123	51.568	-13.390	1.00	110.35
	ATOM	42	N	ALA	7	-35.927	50.819	-11.194	1.00	109.87
	ATOM	43	CA	ALA	7	-34.535	50.380	-11.088	1.00	107.78
45	ATOM	44	C	ALA	7	-33.551	51.344	-11.778	1.00	105.15
	ATOM	45	O	ALA	7	-32.521	51.639	-11.196	1.00	103.42
	ATOM	46	CB	ALA	7	-34.377	48.984	-11.653	1.00	109.08
	ATOM	47	N	GLU	8	-33.840	51.904	-12.996	1.00	101.11
	ATOM	48	CA	GLU	8	-32.804	52.766	-13.697	1.00	95.31
50	ATOM	49	CB	GLU	8	-32.741	52.502	-15.179	1.00	92.47
	ATOM	50	CG	GLU	8	-33.558	51.311	-15.644	1.00	86.82
	ATOM	51	CD	GLU	8	-34.390	51.639	-16.872	1.00	85.55
	ATOM	52	OE1	GLU	8	-35.165	52.622	-16.827	1.00	85.72
	ATOM	53	OE2	GLU	8	-34.254	50.930	-17.879	1.00	81.01
55	ATOM	54	C	GLU	8	-32.884	54.272	-13.535	1.00	92.66
	ATOM	55	O	GLU	8	-33.909	54.825	-13.120	1.00	94.43
	ATOM	56	N	CYS	9	-31.784	54.932	-13.880	1.00	85.19
	ATOM	57	CA	CYS	9	-31.682	56.376	-13.744	1.00	78.14
	ATOM	58	C	CYS	9	-31.252	56.961	-15.074	1.00	75.43
60	ATOM	59	O	CYS	9	-30.372	56.426	-15.745	1.00	76.93
	ATOM	60	CB	CYS	9	-30.675	56.751	-12.658	1.00	76.59
	ATOM	61	SG	CYS	9	-28.942	56.366	-13.062	1.00	20.00
	ATOM	62	N	PHE	10	-31.900	58.068	-15.429	1.00	69.11
	ATOM	63	CA	PHE	10	-31.610	58.835	-16.665	1.00	62.54
65	ATOM	64	CB	PHE	10	-32.665	59.992	-16.855	1.00	61.14
	ATOM	65	CG	PHE	10	-32.511	60.790	-18.130	1.00	61.11

	ATOM	66	CD1	PHE	10	-32.829	60.191	-19.344	1.00	59.90
	ATOM	67	CD2	PHE	10	-32.016	62.093	-18.131	1.00	63.23
	ATOM	68	CE1	PHE	10	-32.654	60.881	-20.544	1.00	61.45
	ATOM	69	CE2	PHE	10	-31.833	62.801	-19.326	1.00	61.15
5	ATOM	70	CZ	PHE	10	-32.151	62.195	-20.534	1.00	61.89
	ATOM	71	C	PHE	10	-30.186	59.413	-16.598	1.00	60.79
	ATOM	72	O	PHE	10	-29.868	60.218	-15.717	1.00	63.10
	ATOM	73	N	ASP	11	-29.308	58.990	-17.517	1.00	57.06
	ATOM	74	CA	ASP	11	-27.930	59.524	-17.554	1.00	52.20
10	ATOM	75	CB	ASP	11	-26.911	58.435	-18.063	1.00	46.52
	ATOM	76	CG	ASP	11	-25.424	58.832	-18.045	1.00	47.00
	ATOM	77	OD1	ASP	11	-25.140	60.034	-18.239	1.00	49.00
	ATOM	78	OD2	ASP	11	-24.575	57.943	-17.828	1.00	48.43
	ATOM	79	C	ASP	11	-27.930	60.733	-18.512	1.00	53.02
15	ATOM	80	O	ASP	11	-28.228	60.557	-19.689	1.00	53.25
	ATOM	81	N	LEU	12	-27.603	61.949	-18.038	1.00	55.34
	ATOM	82	CA	LEU	12	-27.575	63.161	-18.886	1.00	54.13
	ATOM	83	C	LEU	12	-26.381	63.205	-19.856	1.00	54.96
	ATOM	84	O	LEU	12	-26.370	63.941	-20.840	1.00	54.11
20	ATOM	85	CB	LEU	12	-27.632	64.424	-18.001	1.00	54.65
	ATOM	86	CG	LEU	12	-28.727	64.496	-16.922	1.00	20.00
	ATOM	87	CD1	LEU	12	-28.557	65.754	-16.087	1.00	20.00
	ATOM	88	CD2	LEU	12	-30.102	64.451	-17.571	1.00	20.00
	ATOM	89	N	LEU	13	-25.355	62.385	-19.561	1.00	49.64
25	ATOM	90	CA	LEU	13	-24.164	62.298	-20.397	1.00	46.11
	ATOM	91	CB	LEU	13	-22.925	61.954	-19.619	1.00	40.39
	ATOM	92	CG	LEU	13	-21.782	61.808	-20.599	1.00	34.58
	ATOM	93	CD1	LEU	13	-21.093	63.152	-20.812	1.00	33.04
	ATOM	94	CD2	LEU	13	-20.792	60.762	-20.130	1.00	33.82
30	ATOM	95	C	LEU	13	-24.326	61.261	-21.497	1.00	49.82
	ATOM	96	O	LEU	13	-23.436	61.126	-22.334	1.00	50.94
	ATOM	97	N	VAL	14	-25.441	60.527	-21.522	1.00	51.20
	ATOM	98	CA	VAL	14	-25.656	59.691	-22.720	1.00	51.39
	ATOM	99	C	VAL	14	-26.930	60.139	-23.359	1.00	53.87
35	ATOM	100	O	VAL	14	-27.203	59.859	-24.528	1.00	50.27
	ATOM	101	CB	VAL	14	-25.909	58.208	-22.519	1.00	47.82
	ATOM	102	CG1	VAL	14	-26.017	57.505	-23.869	1.00	20.00
	ATOM	103	CG2	VAL	14	-24.822	57.569	-21.662	1.00	20.00
	ATOM	104	N	ARG	15	-27.734	60.822	-22.532	1.00	58.57
40	ATOM	105	CA	ARG	15	-29.009	61.402	-22.880	1.00	64.33
	ATOM	106	C	ARG	15	-30.069	60.325	-23.142	1.00	66.69
	ATOM	107	O	ARG	15	-30.972	60.471	-23.979	1.00	67.11
	ATOM	108	CB	ARG	15	-28.761	62.413	-24.026	1.00	65.68
	ATOM	109	CG	ARG	15	-27.577	63.374	-23.818	1.00	69.20
45	ATOM	110	CD	ARG	15	-27.935	64.835	-24.076	1.00	20.00
	ATOM	111	NE	ARG	15	-26.777	65.761	-24.076	1.00	20.00
	ATOM	112	CZ	ARG	15	-26.874	67.065	-24.307	1.00	20.00
	ATOM	113	NH1	ARG	15	-28.056	67.604	-24.575	1.00	20.00
	ATOM	114	NH2	ARG	15	-25.789	67.826	-24.269	1.00	20.00
50	ATOM	115	N	HIS	16	-29.903	59.229	-22.378	1.00	69.57
	ATOM	116	CA	HIS	16	-30.781	58.072	-22.416	1.00	71.62
	ATOM	117	C	HIS	16	-30.807	57.281	-21.144	1.00	73.99
	ATOM	118	O	HIS	16	-29.917	57.387	-20.310	1.00	75.45
	ATOM	119	CB	HIS	16	-30.330	57.024	-23.401	1.00	69.66
55	ATOM	120	CG	HIS	16	-30.084	57.547	-24.776	1.00	20.00
	ATOM	121	ND1	HIS	16	-31.123	57.701	-25.669	1.00	20.00
	ATOM	122	CD2	HIS	16	-28.951	57.939	-25.412	1.00	20.00
	ATOM	123	CE1	HIS	16	-30.638	58.168	-26.796	1.00	20.00
	ATOM	124	NE2	HIS	16	-29.330	58.310	-26.678	1.00	20.00
60	ATOM	125	N	CYS	17	-31.830	56.468	-21.010	1.00	75.75
	ATOM	126	CA	CYS	17	-31.942	55.693	-19.795	1.00	77.01
	ATOM	127	C	CYS	17	-30.868	54.594	-19.709	1.00	75.86
	ATOM	128	O	CYS	17	-30.424	54.033	-20.708	1.00	73.08
	ATOM	129	CB	CYS	17	-33.344	55.098	-19.688	1.00	80.96
65	ATOM	130	SG	CYS	17	-34.259	55.602	-18.191	1.00	86.58
	ATOM	131	N	VAL	18	-30.468	54.313	-18.469	1.00	75.89

	ATOM	132	CA	VAL	18	-29.480	53.323	-18.148	1.00	77.02
	ATOM	133	C	VAL	18	-29.985	52.520	-16.992	1.00	78.19
	ATOM	134	O	VAL	18	-30.784	52.974	-16.188	1.00	77.54
	ATOM	135	CB	VAL	18	-28.092	53.910	-17.775	1.00	76.68
5	ATOM	136	CG1	VAL	18	-27.404	53.046	-16.740	1.00	76.28
	ATOM	137	CG2	VAL	18	-27.225	54.029	-19.011	1.00	76.37
	ATOM	138	N	ALA	19	-29.461	51.294	-16.909	1.00	80.09
	ATOM	139	CA	ALA	19	-29.781	50.313	-15.821	1.00	82.16
	ATOM	140	C	ALA	19	-28.862	50.487	-14.644	1.00	84.28
10	ATOM	141	O	ALA	19	-27.650	50.523	-14.861	1.00	84.12
	ATOM	142	CB	ALA	19	-29.690	48.882	-16.355	1.00	80.00
	ATOM	143	N	CYS	20	-29.303	50.647	-13.427	1.00	87.37
	ATOM	144	CA	CYS	20	-28.221	50.994	-12.484	1.00	88.33
	ATOM	145	CB	CYS	20	-28.817	51.504	-11.166	1.00	86.90
15	ATOM	146	SG	CYS	20	-28.817	53.316	-10.988	1.00	84.56
	ATOM	147	C	CYS	20	-27.176	49.895	-12.285	1.00	89.64
	ATOM	148	O	CYS	20	-26.219	50.074	-11.538	1.00	89.77
	ATOM	149	N	GLY	21	-27.367	48.751	-12.941	1.00	87.37
	ATOM	150	CA	GLY	21	-26.413	47.679	-12.777	1.00	88.33
20	ATOM	151	C	GLY	21	-25.045	48.073	-13.334	1.00	89.64
	ATOM	152	O	GLY	21	-24.026	47.425	-13.104	1.00	89.77
	ATOM	153	N	LEU	22	-25.020	49.163	-14.109	1.00	87.56
	ATOM	154	CA	LEU	22	-23.799	49.657	-14.748	1.00	85.33
	ATOM	155	CB	LEU	22	-24.124	50.440	-16.007	1.00	79.98
25	ATOM	156	CG	LEU	22	-23.676	49.806	-17.315	1.00	77.91
	ATOM	157	CD1	LEU	22	-24.371	50.451	-18.499	1.00	78.13
	ATOM	158	CD2	LEU	22	-22.157	49.899	-17.461	1.00	72.50
	ATOM	159	C	LEU	22	-22.963	50.527	-13.827	1.00	86.05
	ATOM	160	O	LEU	22	-21.763	50.312	-13.726	1.00	88.06
30	ATOM	161	N	LEU	23	-23.594	51.526	-13.159	1.00	85.11
	ATOM	162	CA	LEU	23	-22.814	52.453	-12.332	1.00	83.33
	ATOM	163	C	LEU	23	-22.505	51.973	-10.916	1.00	84.21
	ATOM	164	O	LEU	23	-21.826	52.693	-10.167	1.00	86.28
	ATOM	165	CB	LEU	23	-23.518	53.811	-12.246	1.00	78.93
35	ATOM	166	CG	LEU	23	-24.137	54.455	-13.478	1.00	73.68
	ATOM	167	CD1	LEU	23	-25.124	55.519	-13.052	1.00	20.00
	ATOM	168	CD2	LEU	23	-23.081	55.022	-14.395	1.00	20.00
	ATOM	169	N	ARG	24	-22.948	50.788	-10.529	1.00	86.09
	ATOM	170	CA	ARG	24	-22.659	50.403	-9.144	1.00	85.39
40	ATOM	171	C	ARG	24	-21.399	49.582	-9.000	1.00	84.10
	ATOM	172	O	ARG	24	-20.796	49.573	-7.929	1.00	83.24
	ATOM	173	CB	ARG	24	-23.826	49.704	-8.507	1.00	80.05
	ATOM	174	CG	ARG	24	-24.699	48.907	-9.431	1.00	20.00
	ATOM	175	CD	ARG	24	-25.479	47.921	-8.607	1.00	20.00
45	ATOM	176	NE	ARG	24	-26.855	48.321	-8.472	1.00	20.00
	ATOM	177	CZ	ARG	24	-27.742	47.470	-7.984	1.00	20.00
	ATOM	178	NH1	ARG	24	-27.362	46.258	-7.602	1.00	20.00
	ATOM	179	NH2	ARG	24	-29.004	47.832	-7.875	1.00	20.00
	ATOM	180	N	THR	25	-20.975	48.860	-10.057	1.00	84.61
50	ATOM	181	CA	THR	25	-19.739	48.049	-9.967	1.00	85.07
	ATOM	182	C	THR	25	-18.540	48.991	-10.079	1.00	85.65
	ATOM	183	O	THR	25	-17.472	48.765	-9.523	1.00	86.77
	ATOM	184	CB	THR	25	-19.763	46.926	-11.021	1.00	20.00
	ATOM	185	OG1	THR	25	-21.019	46.253	-10.951	1.00	20.00
55	ATOM	186	CG2	THR	25	-18.630	45.944	-10.773	1.00	20.00
	ATOM	187	N	PRO	26	-18.798	50.087	-10.865	1.00	84.19
	ATOM	188	CA	PRO	26	-17.903	51.314	-10.951	1.00	82.44
	ATOM	189	C	PRO	26	-17.925	52.220	-9.706	1.00	84.32
	ATOM	190	O	PRO	26	-17.116	53.132	-9.524	1.00	81.24
60	ATOM	191	CB	PRO	26	-18.405	52.036	-12.181	1.00	78.00
	ATOM	192	CG	PRO	26	-18.721	50.913	-13.108	1.00	20.00
	ATOM	193	CD	PRO	26	-18.930	49.680	-12.262	1.00	20.00
	ATOM	194	N	ARG	27	-18.927	51.931	-8.839	1.00	87.06
	ATOM	195	CA	ARG	27	-19.360	52.661	-7.616	1.00	88.17
65	ATOM	196	C	ARG	27	-19.486	54.165	-7.995	1.00	88.45
	ATOM	197	O	ARG	27	-19.358	55.058	-7.160	1.00	84.89

	ATOM	198	CB	ARG	27	-18.594	52.176	-6.338	0.00	86.84
	ATOM	199	CG	ARG	27	-19.113	50.783	-5.799	0.00	85.55
	ATOM	200	CD	ARG	27	-19.481	50.617	-4.273	1.00	20.00
	ATOM	201	NE	ARG	27	-19.966	49.264	-3.871	1.00	20.00
5	ATOM	202	CZ	ARG	27	-20.367	48.851	-2.660	1.00	20.00
	ATOM	203	NH1	ARG	27	-20.360	49.702	-1.654	1.00	20.00
	ATOM	204	NH2	ARG	27	-20.752	47.598	-2.475	1.00	20.00
	ATOM	205	N	PRO	28	-19.757	54.338	-9.305	1.00	89.49
	ATOM	206	CA	PRO	28	-20.149	55.689	-9.836	1.00	91.86
10	ATOM	207	C	PRO	28	-21.395	56.243	-9.059	1.00	95.75
	ATOM	208	O	PRO	28	-21.795	57.397	-9.188	1.00	96.59
	ATOM	209	CB	PRO	28	-20.236	55.456	-11.349	0.00	91.23
	ATOM	210	CG	PRO	28	-19.081	54.509	-11.569	1.00	20.00
	ATOM	211	CD	PRO	28	-18.815	53.778	-10.283	1.00	20.00
15	ATOM	212	N	LYS	29	-21.979	55.334	-8.241	1.00	98.41
	ATOM	213	CA	LYS	29	-23.164	55.475	-7.367	1.00	96.39
	ATOM	214	C	LYS	29	-24.220	56.551	-7.589	1.00	96.88
	ATOM	215	O	LYS	29	-23.986	57.727	-7.322	1.00	94.52
	ATOM	216	CB	LYS	29	-22.676	55.570	-5.922	0.00	95.29
20	ATOM	217	CG	LYS	29	-21.224	55.978	-5.790	0.00	92.34
	ATOM	218	CD	LYS	29	-21.098	57.438	-5.403	0.00	89.50
	ATOM	219	CE	LYS	29	-19.773	57.713	-4.702	1.00	20.00
	ATOM	220	NZ	LYS	29	-19.953	58.539	-3.479	1.00	20.00
	ATOM	221	N	PRO	30	-25.411	56.122	-8.142	1.00	97.72
25	ATOM	222	CA	PRO	30	-26.616	57.021	-8.188	1.00	98.94
	ATOM	223	C	PRO	30	-27.231	57.360	-6.826	1.00	100.71
	ATOM	224	O	PRO	30	-27.621	56.450	-6.082	1.00	100.56
	ATOM	225	CB	PRO	30	-27.571	56.322	-9.118	0.00	97.33
	ATOM	226	CG	PRO	30	-26.666	55.748	-10.140	1.00	20.00
30	ATOM	227	CD	PRO	30	-25.292	55.640	-9.521	1.00	20.00
	ATOM	228	N	ALA	31	-27.335	58.647	-6.490	1.00	101.17
	ATOM	229	CA	ALA	31	-27.905	59.043	-5.224	1.00	99.22
	ATOM	230	C	ALA	31	-29.308	59.619	-5.406	1.00	99.01
	ATOM	231	O	ALA	31	-30.083	59.734	-4.454	1.00	102.17
35	ATOM	232	CB	ALA	31	-26.989	60.048	-4.541	1.00	20.00
	ATOM	233	N	GLY	32	-29.604	59.936	-6.653	1.00	95.52
	ATOM	234	CA	GLY	32	-30.919	60.360	-6.981	1.00	91.44
	ATOM	235	C	GLY	32	-31.637	59.131	-7.512	1.00	88.49
	ATOM	236	O	GLY	32	-31.814	58.986	-8.711	1.00	86.70
40	ATOM	237	N	ALA	33	-32.059	58.257	-6.610	1.00	15.00
	ATOM	238	CA	ALA	33	-32.797	57.070	-7.000	1.00	15.00
	ATOM	239	C	ALA	33	-32.002	55.787	-7.024	1.00	15.00
	ATOM	240	O	ALA	33	-32.540	54.704	-6.782	1.00	15.00
	ATOM	241	CB	ALA	33	-33.427	57.296	-8.374	1.00	15.00
45	ATOM	242	N	SER	34	-30.728	55.877	-7.301	1.00	15.00
	ATOM	243	CA	SER	34	-29.937	54.653	-7.444	1.00	15.00
	ATOM	244	C	SER	34	-29.509	54.004	-6.132	1.00	15.00
	ATOM	245	O	SER	34	-29.416	52.779	-6.027	1.00	15.00
	ATOM	246	CB	SER	34	-28.706	54.957	-8.285	1.00	15.00
50	ATOM	247	OG	SER	34	-28.168	53.765	-8.832	1.00	20.00
	ATOM	248	N	SER	35	-29.241	54.809	-5.121	1.00	15.00
	ATOM	249	CA	SER	35	-28.779	54.302	-3.823	1.00	15.00
	ATOM	250	C	SER	35	-29.616	53.090	-3.413	1.00	15.00
	ATOM	251	O	SER	35	-29.133	52.097	-2.871	1.00	15.00
55	ATOM	252	CB	SER	35	-28.847	55.398	-2.746	1.00	15.00
	ATOM	253	OG	SER	35	-29.130	56.663	-3.326	1.00	20.00
	ATOM	254	N	PRO	36	-30.902	53.244	-3.713	1.00	15.00
	ATOM	255	CA	PRO	36	-31.915	52.184	-3.463	1.00	15.00
	ATOM	256	C	PRO	36	-31.504	50.829	-3.972	1.00	15.00
60	ATOM	257	O	PRO	36	-31.487	49.844	-3.233	1.00	15.00
	ATOM	258	CB	PRO	36	-33.198	52.843	-3.970	1.00	15.00
	ATOM	259	CG	PRO	36	-33.000	54.280	-3.558	1.00	15.00
	ATOM	260	CD	PRO	36	-31.538	54.547	-3.476	1.00	15.00
	ATOM	261	N	ALA	37	-31.156	50.796	-5.245	1.00	15.00
65	ATOM	262	CA	ALA	37	-30.674	49.591	-5.885	1.00	15.00
	ATOM	263	C	ALA	37	-29.387	49.142	-5.212	1.00	15.00

	ATOM	264	O	ALA	37	-29.134	47.953	-5.068	1.00	15.00
	ATOM	265	CB	ALA	37	-30.448	49.834	-7.364	1.00	15.00
	ATOM	266	N	PRO	38	-28.540	50.129	-4.831	1.00	15.00
	ATOM	267	CA	PRO	38	-27.236	49.805	-4.165	1.00	15.00
5	ATOM	268	C	PRO	38	-27.307	48.885	-3.008	1.00	15.00
	ATOM	269	O	PRO	38	-26.627	47.872	-2.929	1.00	15.00
	ATOM	270	CB	PRO	38	-26.629	51.137	-3.854	1.00	15.00
	ATOM	271	CG	PRO	38	-27.000	51.896	-5.095	1.00	20.00
	ATOM	272	CD	PRO	38	-28.247	51.277	-5.685	1.00	20.00
10	ATOM	273	N	ARG	39	-28.186	49.267	-2.092	1.00	15.00
	ATOM	274	CA	ARG	39	-28.384	48.493	-0.868	1.00	15.00
	ATOM	275	C	ARG	39	-29.320	47.287	-1.105	1.00	15.00
	ATOM	276	O	ARG	39	-29.584	46.535	-0.163	1.00	15.00
	ATOM	277	CB	ARG	39	-28.937	49.351	0.247	1.00	15.00
15	ATOM	278	CG	ARG	39	-28.157	50.610	0.506	1.00	15.00
	ATOM	279	CD	ARG	39	-29.130	51.730	0.797	1.00	15.00
	ATOM	280	NE	ARG	39	-28.451	52.972	1.096	1.00	20.00
	ATOM	281	CZ	ARG	39	-28.927	53.885	1.936	1.00	20.00
	ATOM	282	NH1	ARG	39	-30.071	53.668	2.571	1.00	20.00
20	ATOM	283	NH2	ARG	39	-28.259	55.011	2.141	1.00	20.00
	ATOM	284	N	THR	40	-29.832	47.138	-2.329	1.00	15.00
	ATOM	285	CA	THR	40	-30.769	46.066	-2.756	1.00	15.00
	ATOM	286	C	THR	40	-29.968	44.833	-3.187	1.00	15.00
	ATOM	287	O	THR	40	-30.199	43.697	-2.787	1.00	15.00
25	ATOM	288	CB	THR	40	-31.615	46.509	-3.926	1.00	15.00
	ATOM	289	OG1	THR	40	-32.108	47.831	-3.650	1.00	20.00
	ATOM	290	CG2	THR	40	-32.778	45.560	-4.136	1.00	20.00
	ATOM	291	N	ALA	41	-28.992	45.142	-4.036	1.00	15.00
	ATOM	292	CA	ALA	41	-28.085	44.130	-4.500	1.00	15.00
30	ATOM	293	C	ALA	41	-27.013	43.858	-3.448	1.00	15.00
	ATOM	294	O	ALA	41	-26.335	42.848	-3.520	1.00	15.00
	ATOM	295	CB	ALA	41	-27.422	44.536	-5.812	1.00	15.00
	ATOM	296	N	LEU	42	-26.847	44.740	-2.502	1.00	15.00
	ATOM	297	CA	LEU	42	-25.866	44.555	-1.440	1.00	15.00
35	ATOM	298	C	LEU	42	-26.426	43.599	-0.399	1.00	15.00
	ATOM	299	O	LEU	42	-25.746	42.667	0.044	1.00	15.00
	ATOM	300	CB	LEU	42	-25.518	45.879	-0.779	1.00	15.00
	ATOM	301	CG	LEU	42	-24.745	46.891	-1.626	1.00	20.00
	ATOM	302	CD1	LEU	42	-24.447	48.150	-0.818	1.00	20.00
40	ATOM	303	CD2	LEU	42	-23.466	46.280	-2.157	1.00	20.00
	ATOM	304	N	GLN	43	-27.672	43.816	-0.009	1.00	15.00
	ATOM	305	CA	GLN	43	-28.379	42.944	0.943	1.00	15.00
	ATOM	306	C	GLN	43	-28.419	41.527	0.422	1.00	15.00
	ATOM	307	O	GLN	43	-27.869	40.609	1.025	1.00	15.00
45	ATOM	308	CB	GLN	43	-29.822	43.430	1.184	1.00	15.00
	ATOM	309	CG	GLN	43	-29.998	44.906	1.453	1.00	20.00
	ATOM	310	CD	GLN	43	-31.473	45.234	1.567	1.00	20.00
	ATOM	311	OE1	GLN	43	-32.339	44.393	1.344	1.00	20.00
	ATOM	312	NE2	GLN	43	-31.975	46.412	1.927	1.00	20.00
50	END					-119.496	62.638	-15.481	0.00	0.00

TABLE 26

	ATOM	2	CA	PRO	1	52.437	83.641	-2.947	1.00	98.53
	ATOM	3	C	PRO	1	53.020	84.827	-2.281	1.00	101.54
5	ATOM	4	O	PRO	1	54.030	85.317	-2.806	1.00	102.94
	ATOM	5	CB	PRO	1	53.420	82.988	-3.899	1.00	96.91
	ATOM	6	CG	PRO	1	52.493	82.147	-4.700	1.00	20.00
	ATOM	7	CD	PRO	1	51.086	82.709	-4.577	1.00	20.00
	ATOM	8	N	THR	2	52.541	85.378	-1.225	1.00	104.82
10	ATOM	9	CA	THR	2	53.418	86.441	-0.853	1.00	107.58
	ATOM	10	C	THR	2	54.194	85.970	0.365	1.00	108.57
	ATOM	11	O	THR	2	54.375	84.759	0.501	1.00	109.82
	ATOM	12	CB	THR	2	52.667	87.751	-0.700	1.00	20.00
	ATOM	13	OG1	THR	2	52.589	88.413	-1.982	1.00	20.00
15	ATOM	14	CG2	THR	2	53.369	88.641	0.313	1.00	20.00
	ATOM	15	N	PRO	3	54.693	86.829	1.284	1.00	108.52
	ATOM	16	CA	PRO	3	55.592	86.236	2.311	1.00	109.56
	ATOM	17	C	PRO	3	55.158	84.998	3.022	1.00	110.79
	ATOM	18	O	PRO	3	54.074	84.989	3.604	1.00	114.50
20	ATOM	19	CB	PRO	3	55.910	87.401	3.226	1.00	108.23
	ATOM	20	CG	PRO	3	56.047	88.500	2.229	1.00	109.80
	ATOM	21	CD	PRO	3	55.250	88.143	0.990	1.00	110.35
	ATOM	22	C	CYS	4	56.017	81.958	4.628	1.00	101.54
	ATOM	23	O	CYS	4	56.711	81.076	4.106	1.00	102.94
25	ATOM	24	CB	CYS	4	54.143	82.340	3.012	1.00	96.91
	ATOM	25	SG	CYS	4	52.457	82.704	3.596	1.00	92.39
	ATOM	26	N	CYS	4	55.936	83.940	3.062	1.00	93.44
	ATOM	27	CA	CYS	4	55.197	82.963	3.853	1.00	98.53
	ATOM	28	N	VAL	5	55.840	82.147	5.913	1.00	104.82
30	ATOM	29	CA	VAL	5	56.500	81.403	6.956	1.00	107.58
	ATOM	30	C	VAL	5	56.193	79.879	6.902	1.00	108.57
	ATOM	31	O	VAL	5	55.057	79.484	6.621	1.00	109.82
	ATOM	32	CB	VAL	5	56.189	82.164	8.253	1.00	109.37
	ATOM	33	CG1	VAL	5	57.102	81.694	9.377	1.00	20.00
35	ATOM	34	CG2	VAL	5	56.302	83.665	8.054	1.00	20.00
	ATOM	35	N	PRO	6	57.225	79.048	7.162	1.00	108.52
	ATOM	36	CA	PRO	6	56.955	77.595	7.061	1.00	109.56
	ATOM	37	C	PRO	6	55.686	77.110	7.777	1.00	110.79
	ATOM	38	O	PRO	6	55.413	77.535	8.891	1.00	114.50
40	ATOM	39	CB	PRO	6	58.291	76.971	7.438	1.00	108.23
	ATOM	40	CG	PRO	6	59.276	77.887	6.811	1.00	109.80
	ATOM	41	CD	PRO	6	58.608	79.237	6.665	1.00	110.35
	ATOM	42	N	ALA	7	54.921	76.212	7.162	1.00	109.87
	ATOM	43	CA	ALA	7	53.650	75.735	7.707	1.00	107.78
45	ATOM	44	C	ALA	7	52.435	76.497	7.143	1.00	105.15
	ATOM	45	O	ALA	7	51.454	75.858	6.802	1.00	103.42
	ATOM	46	CB	ALA	7	53.659	75.818	9.219	1.00	109.08
	ATOM	47	N	GLU	8	52.464	77.858	6.976	1.00	101.11
	ATOM	48	CA	GLU	8	51.215	78.591	6.518	1.00	95.31
50	ATOM	49	CB	GLU	8	51.006	79.890	7.252	1.00	92.47
	ATOM	50	CG	GLU	8	51.945	80.112	8.422	1.00	86.82
	ATOM	51	CD	GLU	8	52.543	81.509	8.411	1.00	85.55
	ATOM	52	OE1	GLU	8	53.140	81.899	7.381	1.00	85.72
	ATOM	53	OE2	GLU	8	52.400	82.217	9.418	1.00	81.01
55	ATOM	54	C	GLU	8	51.059	78.905	5.043	1.00	92.66
	ATOM	55	O	GLU	8	52.020	78.863	4.266	1.00	94.43
	ATOM	56	N	CYS	9	49.827	79.232	4.666	1.00	85.19
	ATOM	57	CA	CYS	9	49.501	79.520	3.278	1.00	78.14
	ATOM	58	C	CYS	9	48.809	80.866	3.211	1.00	75.43
60	ATOM	59	O	CYS	9	47.953	81.179	4.035	1.00	76.93
	ATOM	60	CB	CYS	9	48.595	78.436	2.696	1.00	76.59
	ATOM	61	SG	CYS	9	46.914	78.394	3.396	1.00	20.00
	ATOM	62	N	PHE	10	49.209	81.646	2.208	1.00	69.11
	ATOM	63	CA	PHE	10	48.635	82.983	1.923	1.00	62.54
65	ATOM	64	CB	PHE	10	49.445	83.695	0.771	1.00	61.14
	ATOM	65	CG	PHE	10	48.993	85.102	0.452	1.00	61.11

	ATOM	66	CD1	PHE	10	49.247	86.116	1.369	1.00	59.90
	ATOM	67	CD2	PHE	10	48.288	85.407	-0.711	1.00	63.23
	ATOM	68	CE1	PHE	10	48.803	87.418	1.130	1.00	61.45
	ATOM	69	CE2	PHE	10	47.835	86.707	-0.967	1.00	61.15
5	ATOM	70	CZ	PHE	10	48.091	87.713	-0.047	1.00	61.89
	ATOM	71	C	PHE	10	47.154	82.843	1.529	1.00	60.79
	ATOM	72	O	PHE	10	46.821	82.201	0.527	1.00	63.10
	ATOM	73	N	ASP	11	46.249	83.422	2.328	1.00	57.06
	ATOM	74	CA	ASP	11	44.807	83.373	2.005	1.00	52.20
10	ATOM	75	CB	ASP	11	43.930	83.344	3.313	1.00	46.52
	ATOM	76	CG	ASP	11	42.412	83.183	3.113	1.00	47.00
	ATOM	77	OD1	ASP	11	41.906	83.676	2.082	1.00	49.00
	ATOM	78	OD2	ASP	11	41.762	82.563	3.980	1.00	48.43
	ATOM	79	C	ASP	11	44.478	84.634	1.180	1.00	53.02
15	ATOM	80	O	ASP	11	44.646	85.737	1.689	1.00	53.25
	ATOM	81	N	LEU	12	44.014	84.496	-0.076	1.00	55.34
	ATOM	82	CA	LEU	12	43.671	85.651	-0.937	1.00	54.13
	ATOM	83	C	LEU	12	42.371	86.362	-0.522	1.00	54.96
	ATOM	84	O	LEU	12	42.108	87.503	-0.893	1.00	54.11
20	ATOM	85	CB	LEU	12	43.627	85.210	-2.415	1.00	54.65
	ATOM	86	CG	LEU	12	44.825	84.414	-2.962	1.00	20.00
	ATOM	87	CD1	LEU	12	44.554	83.978	-4.393	1.00	20.00
	ATOM	88	CD2	LEU	12	46.092	85.252	-2.876	1.00	20.00
	ATOM	89	N	LEU	13	41.546	85.658	0.277	1.00	49.64
25	ATOM	90	CA	LEU	13	40.288	86.206	0.771	1.00	46.11
	ATOM	91	CB	LEU	13	39.237	85.154	0.991	1.00	40.39
	ATOM	92	CG	LEU	13	38.018	85.827	1.582	1.00	34.58
	ATOM	93	CD1	LEU	13	37.090	86.309	0.472	1.00	33.04
	ATOM	94	CD2	LEU	13	37.289	84.898	2.531	1.00	33.82
30	ATOM	95	C	LEU	13	40.480	86.954	2.080	1.00	49.82
	ATOM	96	O	LEU	13	39.525	87.541	2.584	1.00	50.94
	ATOM	97	N	VAL	14	41.691	86.954	2.644	1.00	51.20
	ATOM	98	CA	VAL	14	41.886	87.864	3.790	1.00	51.39
	ATOM	99	C	VAL	14	42.973	88.823	3.423	1.00	53.87
35	ATOM	100	O	VAL	14	43.134	89.883	4.031	1.00	50.27
	ATOM	101	CB	VAL	14	42.412	87.275	5.087	1.00	47.82
	ATOM	102	CG1	VAL	14	42.461	88.349	6.169	1.00	20.00
	ATOM	103	CG2	VAL	14	41.569	86.089	5.540	1.00	20.00
	ATOM	104	N	ARG	15	43.749	88.395	2.418	1.00	58.57
40	ATOM	105	CA	ARG	15	44.851	89.120	1.833	1.00	64.33
	ATOM	106	C	ARG	15	46.036	89.230	2.801	1.00	66.69
	ATOM	107	O	ARG	15	46.785	90.218	2.826	1.00	67.11
	ATOM	108	CB	ARG	15	44.289	90.454	1.284	1.00	65.68
	ATOM	109	CG	ARG	15	42.996	90.338	0.457	1.00	69.20
45	ATOM	110	CD	ARG	15	43.065	91.082	-0.874	1.00	20.00
	ATOM	111	NE	ARG	15	41.777	91.155	-1.604	1.00	20.00
	ATOM	112	CZ	ARG	15	41.620	91.780	-2.766	1.00	20.00
	ATOM	113	NH1	ARG	15	42.648	92.402	-3.327	1.00	20.00
	ATOM	114	NH2	ARG	15	40.436	91.781	-3.363	1.00	20.00
50	ATOM	115	N	HIS	16	46.158	88.155	3.604	1.00	69.57
	ATOM	116	CA	HIS	16	47.207	87.998	4.597	1.00	71.62
	ATOM	117	C	HIS	16	47.532	86.573	4.925	1.00	73.99
	ATOM	118	O	HIS	16	46.752	85.667	4.663	1.00	75.45
	ATOM	119	CB	HIS	16	46.816	88.528	5.952	1.00	69.66
55	ATOM	120	CG	HIS	16	46.309	89.930	5.936	1.00	20.00
	ATOM	121	ND1	HIS	16	47.182	90.998	5.955	1.00	20.00
	ATOM	122	CD2	HIS	16	45.053	90.444	5.909	1.00	20.00
	ATOM	123	CE1	HIS	16	46.483	92.108	5.938	1.00	20.00
	ATOM	124	NE2	HIS	16	45.196	91.810	5.925	1.00	20.00
60	ATOM	125	N	CYS	17	48.687	86.385	5.522	1.00	75.75
	ATOM	126	CA	CYS	17	49.085	85.033	5.844	1.00	77.01
	ATOM	127	C	CYS	17	48.233	84.433	6.977	1.00	75.86
	ATOM	128	O	CYS	17	47.765	85.123	7.879	1.00	73.08
	ATOM	129	CB	CYS	17	50.570	85.002	6.200	1.00	80.96
65	ATOM	130	SG	CYS	17	51.572	83.911	5.133	1.00	86.58
	ATOM	131	N	VAL	18	48.052	83.116	6.887	1.00	75.89

	ATOM	132	CA	VAL	18	47.295	82.344	7.832	1.00	77.02
	ATOM	133	C	VAL	18	48.075	81.107	8.152	1.00	78.19
	ATOM	134	O	VAL	18	48.883	80.631	7.370	1.00	77.54
	ATOM	135	CB	VAL	18	45.889	81.925	7.326	1.00	76.68
5	ATOM	136	CG1	VAL	18	45.497	80.573	7.886	1.00	76.28
	ATOM	137	CG2	VAL	18	44.860	82.965	7.718	1.00	76.37
	ATOM	138	N	ALA	19	47.781	80.570	9.338	1.00	80.09
	ATOM	139	CA	ALA	19	48.402	79.312	9.870	1.00	82.16
	ATOM	140	C	ALA	19	47.628	78.099	9.436	1.00	84.28
10	ATOM	141	O	ALA	19	46.409	78.099	9.618	1.00	84.12
	ATOM	142	CB	ALA	19	48.486	79.367	11.396	1.00	80.00
	ATOM	143	N	CYS	20	48.190	77.083	8.840	1.00	87.37
	ATOM	144	CA	CYS	20	47.196	76.112	8.339	1.00	88.33
	ATOM	145	CB	CYS	20	47.862	75.135	7.363	1.00	86.90
15	ATOM	146	SG	CYS	20	47.578	75.512	5.604	1.00	84.56
	ATOM	147	C	CYS	20	46.387	75.412	9.432	1.00	89.64
	ATOM	148	O	CYS	20	45.519	74.596	9.137	1.00	89.77
	ATOM	149	N	GLY	21	46.683	75.717	10.694	1.00	87.37
	ATOM	150	CA	GLY	21	45.954	75.073	11.762	1.00	88.33
20	ATOM	151	C	GLY	21	44.478	75.472	11.737	1.00	89.64
	ATOM	152	O	GLY	21	43.623	74.881	12.393	1.00	89.77
	ATOM	153	N	LEU	22	44.168	76.521	10.966	1.00	87.56
	ATOM	154	CA	LEU	22	42.809	77.052	10.856	1.00	85.33
	ATOM	155	CB	LEU	22	42.830	78.525	10.488	1.00	79.98
25	ATOM	156	CG	LEU	22	42.330	79.481	11.561	1.00	77.91
	ATOM	157	CD1	LEU	22	42.746	80.907	11.254	1.00	78.13
	ATOM	158	CD2	LEU	22	40.811	79.378	11.702	1.00	72.50
	ATOM	159	C	LEU	22	41.964	76.302	9.842	1.00	86.05
	ATOM	160	O	LEU	22	40.841	75.930	10.156	1.00	88.06
30	ATOM	161	N	LEU	23	42.498	76.087	8.613	1.00	85.11
	ATOM	162	CA	LEU	23	41.686	75.453	7.569	1.00	83.33
	ATOM	163	C	LEU	23	41.649	73.927	7.599	1.00	84.21
	ATOM	164	O	LEU	23	40.961	73.320	6.762	1.00	86.28
	ATOM	165	CB	LEU	23	42.155	75.904	6.181	1.00	78.93
35	ATOM	166	CG	LEU	23	42.490	77.361	5.901	1.00	73.68
	ATOM	167	CD1	LEU	23	43.330	77.455	4.646	1.00	20.00
	ATOM	168	CD2	LEU	23	41.244	78.205	5.792	1.00	20.00
	ATOM	169	N	ARG	24	42.333	73.286	8.534	1.00	86.09
	ATOM	170	CA	ARG	24	42.296	71.822	8.482	1.00	85.39
40	ATOM	171	C	ARG	24	41.223	71.218	9.357	1.00	84.10
	ATOM	172	O	ARG	24	40.774	70.104	9.092	1.00	83.24
	ATOM	173	CB	ARG	24	43.638	71.220	8.792	1.00	80.05
	ATOM	174	CG	ARG	24	44.506	72.002	9.734	1.00	20.00
	ATOM	175	CD	ARG	24	45.541	71.071	10.300	1.00	20.00
45	ATOM	176	NE	ARG	24	46.835	71.307	9.716	1.00	20.00
	ATOM	177	CZ	ARG	24	47.910	70.751	10.250	1.00	20.00
	ATOM	178	NH1	ARG	24	47.794	69.963	11.311	1.00	20.00
	ATOM	179	NH2	ARG	24	49.095	70.981	9.725	1.00	20.00
	ATOM	180	N	THR	25	40.794	71.917	10.427	1.00	84.61
50	ATOM	181	CA	THR	25	39.736	71.370	11.307	1.00	85.07
	ATOM	182	C	THR	25	38.390	71.547	10.604	1.00	85.65
	ATOM	183	O	THR	25	37.457	70.769	10.764	1.00	86.77
	ATOM	184	CB	THR	25	39.813	72.026	12.699	1.00	20.00
	ATOM	185	OG1	THR	25	41.162	71.980	13.158	1.00	20.00
55	ATOM	186	CG2	THR	25	38.905	71.299	13.677	1.00	20.00
	ATOM	187	N	PRO	26	38.354	72.657	9.798	1.00	84.19
	ATOM	188	CA	PRO	26	37.260	72.948	8.781	1.00	82.44
	ATOM	189	C	PRO	26	37.290	72.057	7.526	1.00	84.32
	ATOM	190	O	PRO	26	36.369	72.018	6.707	1.00	81.24
60	ATOM	191	CB	PRO	26	37.469	74.407	8.440	1.00	78.00
	ATOM	192	CG	PRO	26	37.848	74.995	9.756	1.00	20.00
	ATOM	193	CD	PRO	26	38.370	73.868	10.615	1.00	20.00
	ATOM	194	N	ARG	27	38.430	71.335	7.397	1.00	87.06
	ATOM	195	CA	ARG	27	38.888	70.484	6.264	1.00	88.17
65	ATOM	196	C	ARG	27	38.708	71.313	4.960	1.00	88.45
	ATOM	197	O	ARG	27	38.540	70.775	3.867	1.00	84.89

	ATOM	198	CB	ARG	27	38.388	69.005	6.398	0.00	86.84
	ATOM	199	CG	ARG	27	39.201	68.173	7.470	0.00	85.55
	ATOM	200	CD	ARG	27	39.788	66.758	7.089	1.00	20.00
	ATOM	201	NE	ARG	27	40.543	66.061	8.170	1.00	20.00
5	ATOM	202	CZ	ARG	27	41.162	64.872	8.119	1.00	20.00
	ATOM	203	NH1	ARG	27	41.141	64.183	6.996	1.00	20.00
	ATOM	204	NH2	ARG	27	41.774	64.391	9.190	1.00	20.00
	ATOM	205	N	PRO	28	38.773	72.640	5.188	1.00	89.49
	ATOM	206	CA	PRO	28	38.858	73.613	4.044	1.00	91.86
10	ATOM	207	C	PRO	28	40.082	73.271	3.124	1.00	95.75
	ATOM	208	O	PRO	28	40.261	73.809	2.034	1.00	96.59
	ATOM	209	CB	PRO	28	38.786	74.977	4.742	0.00	91.23
	ATOM	210	CG	PRO	28	37.789	74.694	5.840	1.00	20.00
	ATOM	211	CD	PRO	28	37.821	73.223	6.142	1.00	20.00
15	ATOM	212	N	LYS	29	40.913	72.334	3.642	1.00	98.41
	ATOM	213	CA	LYS	29	42.160	71.766	3.086	1.00	96.39
	ATOM	214	C	LYS	29	42.981	72.485	2.021	1.00	96.88
	ATOM	215	O	LYS	29	42.587	72.545	0.859	1.00	94.52
	ATOM	216	CB	LYS	29	41.855	70.355	2.588	0.00	95.29
20	ATOM	217	CG	LYS	29	40.384	70.097	2.337	0.00	92.34
	ATOM	218	CD	LYS	29	40.064	70.149	0.857	0.00	89.50
	ATOM	219	CE	LYS	29	38.814	69.341	0.531	1.00	20.00
	ATOM	220	NZ	LYS	29	39.008	68.474	-0.661	1.00	20.00
	ATOM	221	N	PRO	30	44.145	73.084	2.460	1.00	97.72
25	ATOM	222	CA	PRO	30	45.163	73.610	1.486	1.00	98.94
	ATOM	223	C	PRO	30	45.884	72.544	0.654	1.00	100.71
	ATOM	224	O	PRO	30	46.516	71.642	1.221	1.00	100.56
	ATOM	225	CB	PRO	30	46.095	74.441	2.328	0.00	97.33
	ATOM	226	CG	PRO	30	45.175	75.067	3.305	1.00	20.00
30	ATOM	227	CD	PRO	30	43.932	74.212	3.371	1.00	20.00
	ATOM	228	N	ALA	31	45.811	72.633	-0.676	1.00	101.17
	ATOM	229	CA	ALA	31	46.465	71.666	-1.525	1.00	99.22
	ATOM	230	C	ALA	31	47.715	72.257	-2.175	1.00	99.01
	ATOM	231	O	ALA	31	48.576	71.537	-2.684	1.00	102.17
35	ATOM	232	CB	ALA	31	45.489	71.166	-2.579	1.00	20.00
	ATOM	233	N	GLY	32	47.788	73.573	-2.105	1.00	95.52
	ATOM	234	CA	GLY	32	48.958	74.241	-2.555	1.00	91.44
	ATOM	235	C	GLY	32	49.799	74.496	-1.316	1.00	88.49
	ATOM	236	O	GLY	32	49.840	75.609	-0.814	1.00	86.70
40	ATOM	237	N	ALA	33	50.477	73.463	-0.837	1.00	15.00
	ATOM	238	CA	ALA	33	51.348	73.603	0.315	1.00	15.00
	ATOM	239	C	ALA	33	50.786	73.100	1.622	1.00	15.00
	ATOM	240	O	ALA	33	51.527	72.644	2.496	1.00	15.00
	ATOM	241	CB	ALA	33	51.747	75.070	0.471	1.00	15.00
45	ATOM	242	N	SER	34	49.490	73.162	1.780	1.00	15.00
	ATOM	243	CA	SER	34	48.907	72.789	3.070	1.00	15.00
	ATOM	244	C	SER	34	48.769	71.289	3.307	1.00	15.00
	ATOM	245	O	SER	34	48.899	70.806	4.434	1.00	15.00
	ATOM	246	CB	SER	34	47.543	73.450	3.204	1.00	15.00
50	ATOM	247	OG	SER	34	47.149	73.511	4.564	1.00	20.00
	ATOM	248	N	SER	35	48.501	70.535	2.256	1.00	15.00
	ATOM	249	CA	SER	35	48.305	69.084	2.368	1.00	15.00
	ATOM	250	C	SER	35	49.381	68.485	3.273	1.00	15.00
	ATOM	251	O	SER	35	49.148	67.593	4.088	1.00	15.00
55	ATOM	252	CB	SER	35	48.325	68.415	0.984	1.00	15.00
	ATOM	253	OG	SER	35	48.312	69.389	-0.050	1.00	20.00
	ATOM	254	N	PRO	36	50.572	69.039	3.071	1.00	15.00
	ATOM	255	CA	PRO	36	51.775	68.665	3.864	1.00	15.00
	ATOM	256	C	PRO	36	51.536	68.664	5.350	1.00	15.00
60	ATOM	257	O	PRO	36	51.782	67.672	6.037	1.00	15.00
	ATOM	258	CB	PRO	36	52.850	69.565	3.255	1.00	15.00
	ATOM	259	CG	PRO	36	52.466	69.575	1.797	1.00	15.00
	ATOM	260	CD	PRO	36	51.003	69.320	1.695	1.00	15.00
	ATOM	261	N	ALA	37	51.037	69.786	5.834	1.00	15.00
65	ATOM	262	CA	ALA	37	50.687	69.939	7.230	1.00	15.00
	ATOM	263	C	ALA	37	49.594	68.945	7.588	1.00	15.00

	ATOM	264	O	ALA	37	49.566	68.409	8.688	1.00	15.00
	ATOM	265	CB	ALA	37	50.233	71.358	7.507	1.00	15.00
	ATOM	266	N	PRO	38	48.648	68.734	6.640	1.00	15.00
	ATOM	267	CA	PRO	38	47.516	67.783	6.885	1.00	15.00
5	ATOM	268	C	PRO	38	47.891	66.435	7.366	1.00	15.00
	ATOM	269	O	PRO	38	47.408	65.937	8.374	1.00	15.00
	ATOM	270	CB	PRO	38	46.737	67.784	5.607	1.00	15.00
	ATOM	271	CG	PRO	38	46.809	69.241	5.251	1.00	20.00
	ATOM	272	CD	PRO	38	48.056	69.829	5.873	1.00	20.00
10	ATOM	273	N	ARG	39	48.804	65.846	6.606	1.00	15.00
	ATOM	274	CA	ARG	39	49.287	64.502	6.914	1.00	15.00
	ATOM	275	C	ARG	39	50.375	64.527	8.011	1.00	15.00
	ATOM	276	O	ARG	39	50.884	63.465	8.380	1.00	15.00
	ATOM	277	CB	ARG	39	49.827	63.812	5.682	1.00	15.00
15	ATOM	278	CG	ARG	39	48.885	63.809	4.510	1.00	15.00
	ATOM	279	CD	ARG	39	49.683	64.043	3.248	1.00	15.00
	ATOM	280	NE	ARG	39	48.848	64.016	2.066	1.00	20.00
	ATOM	281	CZ	ARG	39	49.268	63.587	0.881	1.00	20.00
	ATOM	282	NH1	ARG	39	50.505	63.129	0.743	1.00	20.00
20	ATOM	283	NH2	ARG	39	48.451	63.615	-0.162	1.00	20.00
	ATOM	284	N	THR	40	50.742	65.720	8.485	1.00	15.00
	ATOM	285	CA	THR	40	51.783	65.964	9.516	1.00	15.00
	ATOM	286	C	THR	40	51.154	65.857	10.909	1.00	15.00
	ATOM	287	O	THR	40	51.624	65.182	11.818	1.00	15.00
25	ATOM	288	CB	THR	40	52.383	67.344	9.377	1.00	15.00
	ATOM	289	OG1	THR	40	52.677	67.569	7.988	1.00	20.00
	ATOM	290	CG2	THR	40	53.653	67.462	10.196	1.00	20.00
	ATOM	291	N	ALA	41	50.038	66.573	11.010	1.00	15.00
	ATOM	292	CA	ALA	41	49.263	66.544	12.220	1.00	15.00
30	ATOM	293	C	ALA	41	48.398	65.287	12.263	1.00	15.00
	ATOM	294	O	ALA	41	47.898	64.932	13.316	1.00	15.00
	ATOM	295	CB	ALA	41	48.376	67.778	12.341	1.00	15.00
	ATOM	296	N	LEU	42	48.209	64.635	11.150	1.00	15.00
	ATOM	297	CA	LEU	42	47.420	63.411	11.097	1.00	15.00
35	ATOM	298	C	LEU	42	48.265	62.247	11.591	1.00	15.00
	ATOM	299	O	LEU	42	47.817	61.432	12.405	1.00	15.00
	ATOM	300	CB	LEU	42	46.942	63.128	9.683	1.00	15.00
	ATOM	301	CG	LEU	42	45.905	64.088	9.099	1.00	20.00
	ATOM	302	CD1	LEU	42	45.506	63.657	7.692	1.00	20.00
40	ATOM	303	CD2	LEU	42	44.690	64.177	9.998	1.00	20.00
	ATOM	304	N	GLN	43	49.496	62.167	11.112	1.00	15.00
	ATOM	305	CA	GLN	43	50.458	61.138	11.539	1.00	15.00
	ATOM	306	C	GLN	43	50.670	61.208	13.033	1.00	15.00
	ATOM	307	O	GLN	43	50.367	60.270	13.765	1.00	15.00
45	ATOM	308	CB	GLN	43	51.817	61.313	10.831	1.00	15.00
	ATOM	309	CG	GLN	43	51.773	61.535	9.338	1.00	20.00
	ATOM	310	CD	GLN	43	53.174	61.786	8.816	1.00	20.00
	ATOM	311	OE1	GLN	43	54.133	61.896	9.574	1.00	20.00
	ATOM	312	NE2	GLN	43	53.512	61.891	7.534	1.00	20.00
50	END					134.002	98.540	-12.573	0.00	0.00

TABLE 27

	ATOM	1	N	PRO	1	23.561	107.465	26.442	1.00	93.44
	ATOM	2	CA	PRO	1	23.338	107.095	27.840	1.00	98.53
5	ATOM	3	C	PRO	1	21.951	106.684	28.152	1.00	101.54
	ATOM	4	O	PRO	1	21.221	107.542	28.668	1.00	102.94
	ATOM	5	CB	PRO	1	23.738	108.348	28.594	1.00	96.91
	ATOM	6	CG	PRO	1	24.935	108.755	27.813	1.00	20.00
	ATOM	7	CD	PRO	1	24.867	108.114	26.436	1.00	20.00
10	ATOM	8	N	THR	2	21.463	105.521	27.908	1.00	104.82
	ATOM	9	CA	THR	2	20.130	105.493	28.416	1.00	107.58
	ATOM	10	C	THR	2	20.155	104.653	29.682	1.00	108.57
	ATOM	11	O	THR	2	21.207	104.603	30.320	1.00	109.82
	ATOM	12	CB	THR	2	19.139	105.063	27.349	1.00	20.00
15	ATOM	13	OG1	THR	2	18.710	106.220	26.597	1.00	20.00
	ATOM	14	CG2	THR	2	17.945	104.376	27.991	1.00	20.00
	ATOM	15	N	PRO	3	19.069	103.978	30.126	1.00	108.52
	ATOM	16	CA	PRO	3	19.191	103.362	31.475	1.00	109.56
	ATOM	17	C	PRO	3	20.402	102.550	31.791	1.00	110.79
20	ATOM	18	O	PRO	3	20.701	101.608	31.058	1.00	114.50
	ATOM	19	CB	PRO	3	17.881	102.622	31.652	1.00	108.23
	ATOM	20	CG	PRO	3	16.935	103.591	31.029	1.00	109.80
	ATOM	21	CD	PRO	3	17.693	104.448	30.035	1.00	110.35
	ATOM	22	C	CYS	4	22.749	101.400	34.185	1.00	101.54
25	ATOM	23	O	CYS	4	23.405	102.148	34.920	1.00	102.94
	ATOM	24	CB	CYS	4	23.226	102.203	31.861	1.00	96.91
	ATOM	25	SG	CYS	4	23.378	101.033	30.474	1.00	92.39
	ATOM	26	N	CYS	4	21.124	102.809	32.858	1.00	93.44
	ATOM	27	CA	CYS	4	22.184	101.808	32.844	1.00	98.53
30	ATOM	28	N	VAL	5	22.469	100.139	34.411	1.00	104.82
	ATOM	29	CA	VAL	5	22.811	99.421	35.613	1.00	107.58
	ATOM	30	C	VAL	5	24.343	99.368	35.875	1.00	108.57
	ATOM	31	O	VAL	5	25.128	99.211	34.935	1.00	109.82
	ATOM	32	CB	VAL	5	22.040	98.095	35.523	1.00	109.37
35	ATOM	33	CG1	VAL	5	22.030	97.394	36.874	1.00	20.00
	ATOM	34	CG2	VAL	5	20.626	98.311	35.014	1.00	20.00
	ATOM	35	N	PRO	6	24.740	99.516	37.157	1.00	108.52
	ATOM	36	CA	PRO	6	26.199	99.520	37.409	1.00	109.56
	ATOM	37	C	PRO	6	26.987	98.388	36.734	1.00	110.79
40	ATOM	38	O	PRO	6	26.541	97.249	36.733	1.00	114.50
	ATOM	39	CB	PRO	6	26.286	99.670	38.921	1.00	108.23
	ATOM	40	CG	PRO	6	25.180	100.611	39.227	1.00	109.80
	ATOM	41	CD	PRO	6	24.162	100.489	38.114	1.00	110.35
	ATOM	42	N	ALA	7	28.160	98.682	36.179	1.00	109.87
45	ATOM	43	CA	ALA	7	28.963	97.709	35.439	1.00	107.78
	ATOM	44	C	ALA	7	28.730	97.776	33.917	1.00	105.15
	ATOM	45	O	ALA	7	29.699	97.734	33.178	1.00	103.42
	ATOM	46	CB	ALA	7	28.691	96.307	35.944	1.00	109.08
	ATOM	47	N	GLU	8	27.471	97.932	33.396	1.00	101.11
50	ATOM	48	CA	GLU	8	27.264	97.889	31.892	1.00	95.31
	ATOM	49	CB	GLU	8	26.029	97.120	31.501	1.00	92.47
	ATOM	50	CG	GLU	8	25.358	96.380	32.643	1.00	86.82
	ATOM	51	CD	GLU	8	23.855	96.602	32.656	1.00	85.55
	ATOM	52	OE1	GLU	8	23.421	97.777	32.674	1.00	85.72
55	ATOM	53	OE2	GLU	8	23.115	95.608	32.629	1.00	81.01
	ATOM	54	C	GLU	8	27.210	99.199	31.130	1.00	92.66
	ATOM	55	O	GLU	8	27.025	100.277	31.707	1.00	94.43
	ATOM	56	N	CYS	9	27.366	99.090	29.815	1.00	85.19
	ATOM	57	CA	CYS	9	27.382	100.255	28.945	1.00	78.14
60	ATOM	58	C	CYS	9	26.366	100.051	27.840	1.00	75.43
	ATOM	59	O	CYS	9	26.257	98.966	27.274	1.00	76.93
	ATOM	60	CB	CYS	9	28.771	100.468	28.346	1.00	76.59
	ATOM	61	SG	CYS	9	29.286	99.195	27.150	1.00	20.00
	ATOM	62	N	PHE	10	25.631	101.125	27.557	1.00	69.11
65	ATOM	63	CA	PHE	10	24.611	101.167	26.481	1.00	62.54
	ATOM	64	CB	PHE	10	23.821	102.532	26.526	1.00	61.14

	ATOM	65	CG	PHE	10	22.699	102.650	25.519	1.00	61.11
	ATOM	66	CD1	PHE	10	21.551	101.886	25.695	1.00	59.90
	ATOM	67	CD2	PHE	10	22.798	103.467	24.394	1.00	63.23
	ATOM	68	CE1	PHE	10	20.515	101.933	24.761	1.00	61.45
5	ATOM	69	CE2	PHE	10	21.769	103.527	23.446	1.00	61.15
	ATOM	70	CZ	PHE	10	20.627	102.760	23.628	1.00	61.89
	ATOM	71	C	PHE	10	25.289	100.985	25.112	1.00	60.79
	ATOM	72	O	PHE	10	26.126	101.797	24.704	1.00	63.10
	ATOM	73	N	ASP	11	24.951	99.903	24.399	1.00	57.06
10	ATOM	74	CA	ASP	11	25.521	99.669	23.055	1.00	52.20
	ATOM	75	CB	ASP	11	25.677	98.129	22.762	1.00	46.52
	ATOM	76	CG	ASP	11	26.362	97.753	21.436	1.00	47.00
	ATOM	77	OD1	ASP	11	26.202	98.520	20.461	1.00	49.00
	ATOM	78	OD2	ASP	11	27.049	96.712	21.397	1.00	48.43
15	ATOM	79	C	ASP	11	24.559	100.304	22.030	1.00	53.02
	ATOM	80	O	ASP	11	23.408	99.885	21.958	1.00	53.25
	ATOM	81	N	LEU	12	25.002	101.299	21.240	1.00	55.34
	ATOM	82	CA	LEU	12	24.148	101.963	20.229	1.00	54.13
	ATOM	83	C	LEU	12	23.868	101.090	18.993	1.00	54.96
20	ATOM	84	O	LEU	12	22.938	101.329	18.227	1.00	54.11
	ATOM	85	CB	LEU	12	24.761	103.323	19.834	1.00	54.65
	ATOM	86	CG	LEU	12	25.172	104.281	20.966	1.00	20.00
	ATOM	87	CD1	LEU	12	25.851	105.512	20.389	1.00	20.00
	ATOM	88	CD2	LEU	12	23.954	104.664	21.793	1.00	20.00
25	ATOM	89	N	LEU	13	24.701	100.048	18.809	1.00	49.64
	ATOM	90	CA	LEU	13	24.549	99.119	17.696	1.00	46.11
	ATOM	91	CB	LEU	13	25.856	98.532	17.240	1.00	40.39
	ATOM	92	CG	LEU	13	25.562	97.527	16.149	1.00	34.58
	ATOM	93	CD1	LEU	13	25.564	98.210	14.785	1.00	33.04
30	ATOM	94	CD2	LEU	13	26.553	96.382	16.181	1.00	33.82
	ATOM	95	C	LEU	13	23.621	97.968	18.050	1.00	49.82
	ATOM	96	O	LEU	13	23.329	97.142	17.188	1.00	50.94
	ATOM	97	N	VAL	14	23.143	97.894	19.295	1.00	51.20
	ATOM	98	CA	VAL	14	22.083	96.895	19.538	1.00	51.39
35	ATOM	99	C	VAL	14	20.870	97.632	20.007	1.00	53.87
	ATOM	100	O	VAL	14	19.749	97.119	19.979	1.00	50.27
	ATOM	101	CB	VAL	14	22.292	95.890	20.656	1.00	47.82
	ATOM	102	CG1	VAL	14	21.136	94.895	20.691	1.00	20.00
	ATOM	103	CG2	VAL	14	23.625	95.165	20.510	1.00	20.00
40	ATOM	104	N	ARG	15	21.136	98.856	20.483	1.00	58.57
	ATOM	105	CA	ARG	15	20.163	99.803	20.973	1.00	64.33
	ATOM	106	C	ARG	15	19.541	99.342	22.297	1.00	66.69
	ATOM	107	O	ARG	15	18.364	99.589	22.600	1.00	67.11
	ATOM	108	CB	ARG	15	19.176	100.094	19.816	1.00	65.68
45	ATOM	109	CG	ARG	15	19.823	100.384	18.450	1.00	69.20
	ATOM	110	CD	ARG	15	19.274	101.640	17.779	1.00	20.00
	ATOM	111	NE	ARG	15	19.730	101.840	16.383	1.00	20.00
	ATOM	112	CZ	ARG	15	19.346	102.856	15.618	1.00	20.00
	ATOM	113	NH1	ARG	15	18.492	103.754	16.090	1.00	20.00
50	ATOM	114	NH2	ARG	15	19.818	102.971	14.384	1.00	20.00
	ATOM	115	N	HIS	16	20.401	98.650	23.068	1.00	69.57
	ATOM	116	CA	HIS	16	20.070	98.118	24.379	1.00	71.62
	ATOM	117	C	HIS	16	21.250	97.944	25.285	1.00	73.99
	ATOM	118	O	HIS	16	22.391	97.905	24.844	1.00	75.45
55	ATOM	119	CB	HIS	16	19.535	96.710	24.323	1.00	69.66
	ATOM	120	CG	HIS	16	18.399	96.527	23.375	1.00	20.00
	ATOM	121	ND1	HIS	16	17.107	96.825	23.755	1.00	20.00
	ATOM	122	CD2	HIS	16	18.344	96.082	22.094	1.00	20.00
	ATOM	123	CE1	HIS	16	16.307	96.573	22.746	1.00	20.00
60	ATOM	124	NE2	HIS	16	17.019	96.111	21.734	1.00	20.00
	ATOM	125	N	CYS	17	20.963	97.819	26.561	1.00	75.75
	ATOM	126	CA	CYS	17	22.050	97.678	27.503	1.00	77.01
	ATOM	127	C	CYS	17	22.753	96.314	27.375	1.00	75.86
	ATOM	128	O	CYS	17	22.152	95.298	27.037	1.00	73.08
65	ATOM	129	CB	CYS	17	21.536	97.898	28.924	1.00	80.96
	ATOM	130	SG	CYS	17	22.353	99.268	29.811	1.00	86.58

	ATOM	131	N	VAL	18	24.055	96.340	27.659	1.00	75.89
	ATOM	132	CA	VAL	18	24.910	95.188	27.610	1.00	77.02
	ATOM	133	C	VAL	18	25.762	95.189	28.840	1.00	78.19
	ATOM	134	O	VAL	18	26.034	96.218	29.438	1.00	77.54
5	ATOM	135	CB	VAL	18	25.837	95.139	26.366	1.00	76.68
	ATOM	136	CG1	VAL	18	27.159	94.484	26.706	1.00	76.28
	ATOM	137	CG2	VAL	18	25.161	94.386	25.239	1.00	76.37
	ATOM	138	N	ALA	19	26.212	93.982	29.192	1.00	80.09
	ATOM	139	CA	ALA	19	27.111	93.728	30.366	1.00	82.16
10	ATOM	140	C	ALA	19	28.558	93.853	29.979	1.00	84.28
	ATOM	141	O	ALA	19	28.944	93.232	28.987	1.00	84.12
	ATOM	142	CB	ALA	19	26.838	92.341	30.951	1.00	80.00
	ATOM	143	N	CYS	20	29.393	94.622	30.622	1.00	87.37
	ATOM	144	CA	CYS	20	30.696	94.726	29.935	1.00	88.33
15	ATOM	145	CB	CYS	20	31.508	95.887	30.522	1.00	86.90
	ATOM	146	SG	CYS	20	31.474	97.412	29.528	1.00	84.56
	ATOM	147	C	CYS	20	31.483	93.416	29.867	1.00	89.64
	ATOM	148	O	CYS	20	32.573	93.374	29.304	1.00	89.77
	ATOM	149	N	GLY	21	30.939	92.351	30.453	1.00	87.37
20	ATOM	150	CA	GLY	21	31.650	91.094	30.425	1.00	88.33
	ATOM	151	C	GLY	21	31.776	90.567	28.995	1.00	89.64
	ATOM	152	O	GLY	21	32.531	89.645	28.695	1.00	89.77
	ATOM	153	N	LEU	22	30.998	91.160	28.082	1.00	87.56
	ATOM	154	CA	LEU	22	30.972	90.755	26.676	1.00	85.33
25	ATOM	155	CB	LEU	22	29.637	91.093	26.038	1.00	79.98
	ATOM	156	CG	LEU	22	28.776	89.905	25.636	1.00	77.91
	ATOM	157	CD1	LEU	22	27.344	90.333	25.378	1.00	78.13
	ATOM	158	CD2	LEU	22	29.364	89.211	24.407	1.00	72.50
	ATOM	159	C	LEU	22	32.084	91.389	25.860	1.00	86.05
30	ATOM	160	O	LEU	22	32.768	90.684	25.130	1.00	88.06
	ATOM	161	N	LEU	23	32.261	92.730	25.971	1.00	85.11
	ATOM	162	CA	LEU	23	33.258	93.403	25.131	1.00	83.33
	ATOM	163	C	LEU	23	34.691	93.373	25.657	1.00	84.21
	ATOM	164	O	LEU	23	35.594	93.899	24.987	1.00	86.28
35	ATOM	165	CB	LEU	23	32.855	94.862	24.892	1.00	78.93
	ATOM	166	CG	LEU	23	31.418	95.236	24.562	1.00	73.68
	ATOM	167	CD1	LEU	23	31.207	96.712	24.818	1.00	20.00
	ATOM	168	CD2	LEU	23	31.062	94.869	23.142	1.00	20.00
	ATOM	169	N	ARG	24	34.941	92.763	26.805	1.00	86.09
40	ATOM	170	CA	ARG	24	36.327	92.808	27.280	1.00	85.39
	ATOM	171	C	ARG	24	37.140	91.602	26.872	1.00	84.10
	ATOM	172	O	ARG	24	38.364	91.690	26.790	1.00	83.24
	ATOM	173	CB	ARG	24	36.399	93.022	28.765	1.00	80.05
	ATOM	174	CG	ARG	24	35.259	92.463	29.565	1.00	20.00
45	ATOM	175	CD	ARG	24	35.709	92.328	30.993	1.00	20.00
	ATOM	176	NE	ARG	24	35.128	93.349	31.825	1.00	20.00
	ATOM	177	CZ	ARG	24	35.219	93.255	33.141	1.00	20.00
	ATOM	178	NH1	ARG	24	35.860	92.233	33.693	1.00	20.00
	ATOM	179	NH2	ARG	24	34.673	94.181	33.902	1.00	20.00
50	ATOM	180	N	THR	25	36.496	90.444	26.623	1.00	84.61
	ATOM	181	CA	THR	25	37.250	89.238	26.209	1.00	85.07
	ATOM	182	C	THR	25	37.626	89.391	24.735	1.00	85.65
	ATOM	183	O	THR	25	38.645	88.902	24.262	1.00	86.77
	ATOM	184	CB	THR	25	36.436	87.969	26.526	1.00	20.00
55	ATOM	185	OG1	THR	25	35.968	88.044	27.871	1.00	20.00
	ATOM	186	CG2	THR	25	37.296	86.730	26.341	1.00	20.00
	ATOM	187	N	PRO	26	36.703	90.118	24.025	1.00	84.19
	ATOM	188	CA	PRO	26	36.927	90.655	22.619	1.00	82.44
	ATOM	189	C	PRO	26	37.908	91.838	22.527	1.00	84.32
60	ATOM	190	O	PRO	26	38.357	92.257	21.458	1.00	81.24
	ATOM	191	CB	PRO	26	35.539	91.038	22.158	1.00	78.00
	ATOM	192	CG	PRO	26	34.696	89.952	22.734	1.00	20.00
	ATOM	193	CD	PRO	26	35.464	89.356	23.890	1.00	20.00
	ATOM	194	N	ARG	27	38.216	92.386	23.728	1.00	87.06
65	ATOM	195	CA	ARG	27	39.000	93.614	24.035	1.00	88.17
	ATOM	196	C	ARG	27	38.452	94.752	23.127	1.00	88.45

	ATOM	197	O	ARG	27	39.148	95.709	22.793	1.00	84.89
	ATOM	198	CB	ARG	27	40.531	93.315	24.179	0.00	86.84
	ATOM	199	CG	ARG	27	40.899	92.627	25.555	0.00	85.55
	ATOM	200	CD	ARG	27	42.072	93.209	26.437	1.00	20.00
5	ATOM	201	NE	ARG	27	42.332	92.490	27.718	1.00	20.00
	ATOM	202	CZ	ARG	27	43.241	92.776	28.662	1.00	20.00
	ATOM	203	NH1	ARG	27	44.033	93.816	28.500	1.00	20.00
	ATOM	204	NH2	ARG	27	43.349	92.012	29.738	1.00	20.00
	ATOM	205	N	PRO	28	37.162	94.555	22.787	1.00	89.49
10	ATOM	206	CA	PRO	28	36.370	95.641	22.113	1.00	91.86
	ATOM	207	C	PRO	28	36.395	96.952	22.975	1.00	95.75
	ATOM	208	O	PRO	28	35.971	98.027	22.557	1.00	96.59
	ATOM	209	CB	PRO	28	35.033	94.957	21.805	0.00	91.23
	ATOM	210	CG	PRO	28	35.492	93.569	21.426	1.00	20.00
15	ATOM	211	CD	PRO	28	36.817	93.311	22.086	1.00	20.00
	ATOM	212	N	LYS	29	36.925	96.786	24.211	1.00	98.41
	ATOM	213	CA	LYS	29	37.107	97.769	25.301	1.00	96.39
	ATOM	214	C	LYS	29	36.296	99.057	25.383	1.00	96.88
	ATOM	215	O	LYS	29	36.519	99.990	24.615	1.00	94.52
20	ATOM	216	CB	LYS	29	38.590	98.129	25.369	0.00	95.29
	ATOM	217	CG	LYS	29	39.356	97.819	24.100	0.00	92.34
	ATOM	218	CD	LYS	29	39.602	99.074	23.288	0.00	89.50
	ATOM	219	CE	LYS	29	40.817	98.919	22.382	1.00	20.00
	ATOM	220	NZ	LYS	29	41.712	100.105	22.445	1.00	20.00
25	ATOM	221	N	PRO	30	35.290	99.077	26.329	1.00	97.72
	ATOM	222	CA	PRO	30	34.581	100.355	26.683	1.00	98.94
	ATOM	223	C	PRO	30	35.440	101.403	27.399	1.00	100.71
	ATOM	224	O	PRO	30	35.998	101.117	28.467	1.00	100.56
	ATOM	225	CB	PRO	30	33.386	99.912	27.485	0.00	97.33
30	ATOM	226	CG	PRO	30	32.987	98.659	26.805	1.00	20.00
	ATOM	227	CD	PRO	30	34.194	98.143	26.058	1.00	20.00
	ATOM	228	N	ALA	31	35.549	102.610	26.840	1.00	101.17
	ATOM	229	CA	ALA	31	36.340	103.648	27.458	1.00	99.22
	ATOM	230	C	ALA	31	35.451	104.711	28.101	1.00	99.01
35	ATOM	231	O	ALA	31	35.899	105.508	28.928	1.00	102.17
	ATOM	232	CB	ALA	31	37.268	104.269	26.424	1.00	20.00
	ATOM	233	N	GLY	32	34.189	104.663	27.717	1.00	95.52
	ATOM	234	CA	GLY	32	33.230	105.510	28.334	1.00	91.44
	ATOM	235	C	GLY	32	32.553	104.669	29.403	1.00	88.49
40	ATOM	236	O	GLY	32	31.437	104.210	29.215	1.00	86.70
	ATOM	237	N	ALA	33	33.229	104.484	30.527	1.00	15.00
	ATOM	238	CA	ALA	33	32.661	103.736	31.633	1.00	15.00
	ATOM	239	C	ALA	33	33.153	102.317	31.782	1.00	15.00
	ATOM	240	O	ALA	33	33.220	101.784	32.892	1.00	15.00
45	ATOM	241	CB	ALA	33	31.138	103.728	31.507	1.00	15.00
	ATOM	242	N	SER	34	33.510	101.690	30.693	1.00	15.00
	ATOM	243	CA	SER	34	33.891	100.278	30.771	1.00	15.00
	ATOM	244	C	SER	34	35.308	100.018	31.272	1.00	15.00
	ATOM	245	O	SER	34	35.572	99.023	31.951	1.00	15.00
50	ATOM	246	CB	SER	34	33.714	99.643	29.400	1.00	15.00
	ATOM	247	OG	SER	34	33.618	98.233	29.511	1.00	20.00
	ATOM	248	N	SER	35	36.234	100.900	30.944	1.00	15.00
	ATOM	249	CA	SER	35	37.640	100.734	31.333	1.00	15.00
	ATOM	250	C	SER	35	37.724	100.297	32.796	1.00	15.00
55	ATOM	251	O	SER	35	38.532	99.460	33.198	1.00	15.00
	ATOM	252	CB	SER	35	38.433	102.032	31.108	1.00	15.00
	ATOM	253	OG	SER	35	37.659	102.980	30.387	1.00	20.00
	ATOM	254	N	PRO	36	36.833	100.922	33.558	1.00	15.00
	ATOM	255	CA	PRO	36	36.678	100.634	35.010	1.00	15.00
60	ATOM	256	C	PRO	36	36.571	99.166	35.322	1.00	15.00
	ATOM	257	O	PRO	36	37.327	98.626	36.130	1.00	15.00
	ATOM	258	CB	PRO	36	35.554	101.592	35.406	1.00	15.00
	ATOM	259	CG	PRO	36	35.858	102.804	34.562	1.00	15.00
	ATOM	260	CD	PRO	36	36.600	102.358	33.351	1.00	15.00
65	ATOM	261	N	ALA	37	35.630	98.523	34.656	1.00	15.00
	ATOM	262	CA	ALA	37	35.428	97.096	34.785	1.00	15.00

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	ATOM	263	C	ALA	37	36.677	96.365	34.319	1.00	15.00
	ATOM	264	O	ALA	37	37.047	95.337	34.871	1.00	15.00
	ATOM	265	CB	ALA	37	34.220	96.660	33.979	1.00	15.00
	ATOM	266	N	PRO	38	37.311	96.896	33.245	1.00	15.00
5	ATOM	267	CA	PRO	38	38.548	96.256	32.691	1.00	15.00
	ATOM	268	C	PRO	38	39.620	95.958	33.667	1.00	15.00
	ATOM	269	O	PRO	38	40.119	94.847	33.779	1.00	15.00
	ATOM	270	CB	PRO	38	38.970	97.154	31.571	1.00	15.00
	ATOM	271	CG	PRO	38	37.631	97.501	30.987	1.00	20.00
10	ATOM	272	CD	PRO	38	36.585	97.381	32.073	1.00	20.00
	ATOM	273	N	ARG	39	39.959	97.007	34.404	1.00	15.00
	ATOM	274	CA	ARG	39	41.013	96.910	35.412	1.00	15.00
	ATOM	275	C	ARG	39	40.486	96.295	36.727	1.00	15.00
	ATOM	276	O	ARG	39	41.260	96.148	37.677	1.00	15.00
15	ATOM	277	CB	ARG	39	41.632	98.260	35.697	1.00	15.00
	ATOM	278	CG	ARG	39	42.099	98.999	34.474	1.00	15.00
	ATOM	279	CD	ARG	39	41.772	100.465	34.641	1.00	15.00
	ATOM	280	NE	ARG	39	42.227	101.253	33.516	1.00	20.00
	ATOM	281	CZ	ARG	39	42.637	102.513	33.620	1.00	20.00
20	ATOM	282	NH1	ARG	39	42.667	103.103	34.807	1.00	20.00
	ATOM	283	NH2	ARG	39	43.017	103.178	32.539	1.00	20.00
	ATOM	284	N	THR	40	39.189	95.982	36.782	1.00	15.00
	ATOM	285	CA	THR	40	38.481	95.408	37.956	1.00	15.00
	ATOM	286	C	THR	40	38.616	93.882	37.935	1.00	15.00
25	ATOM	287	O	THR	40	38.973	93.217	38.901	1.00	15.00
	ATOM	288	CB	THR	40	37.010	95.750	37.932	1.00	15.00
	ATOM	289	OG1	THR	40	36.877	97.147	37.622	1.00	20.00
	ATOM	290	CG2	THR	40	36.370	95.459	39.275	1.00	20.00
	ATOM	291	N	ALA	41	38.309	93.369	36.747	1.00	15.00
30	ATOM	292	CA	ALA	41	38.443	91.959	36.508	1.00	15.00
	ATOM	293	C	ALA	41	39.901	91.607	36.225	1.00	15.00
	ATOM	294	O	ALA	41	40.267	90.447	36.286	1.00	15.00
	ATOM	295	CB	ALA	41	37.573	91.508	35.340	1.00	15.00
	ATOM	296	N	LEU	42	40.714	92.576	35.907	1.00	15.00
35	ATOM	297	CA	LEU	42	42.128	92.341	35.644	1.00	15.00
	ATOM	298	C	LEU	42	42.869	92.204	36.965	1.00	15.00
	ATOM	299	O	LEU	42	43.677	91.288	37.153	1.00	15.00
	ATOM	300	CB	LEU	42	42.732	93.479	34.839	1.00	15.00
	ATOM	301	CG	LEU	42	42.257	93.630	33.393	1.00	20.00
40	ATOM	302	CD1	LEU	42	42.972	94.792	32.711	1.00	20.00
	ATOM	303	CD2	LEU	42	42.468	92.344	32.622	1.00	20.00
	ATOM	304	N	GLN	43	42.591	93.106	37.893	1.00	15.00
	ATOM	305	CA	GLN	43	43.175	93.074	39.244	1.00	15.00
	ATOM	306	C	GLN	43	42.849	91.765	39.923	1.00	15.00
45	ATOM	307	O	GLN	43	43.734	90.979	40.252	1.00	15.00
	ATOM	308	CB	GLN	43	42.645	94.234	40.111	1.00	15.00
	ATOM	309	CG	GLN	43	42.642	95.603	39.473	1.00	20.00
	ATOM	310	CD	GLN	43	42.003	96.605	40.414	1.00	20.00
	ATOM	311	OE1	GLN	43	41.483	96.256	41.470	1.00	20.00
50	ATOM	312	NE2	GLN	43	41.954	97.920	40.222	1.00	20.00
	END					-16.719	146.167	89.779	0.00	0.00

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	ATOM	2	CA	PRO	1	7.309	67.626	4.820	1.00	98.53
	ATOM	3	C	PRO	1	8.128	66.826	3.882	1.00	101.54
5	ATOM	4	O	PRO	1	7.515	66.067	3.119	1.00	102.94
	ATOM	5	CB	PRO	1	5.910	67.061	4.970	1.00	96.91
	ATOM	6	CG	PRO	1	5.219	68.227	5.581	1.00	20.00
	ATOM	7	CD	PRO	1	6.013	69.487	5.276	1.00	20.00
	ATOM	8	N	THR	2	9.409	66.883	3.802	1.00	104.82
10	ATOM	9	CA	THR	2	9.828	65.917	2.839	1.00	107.58
	ATOM	10	C	THR	2	10.441	64.760	3.610	1.00	108.57
	ATOM	11	O	THR	2	10.045	64.558	4.759	1.00	109.82
	ATOM	12	CB	THR	2	10.704	66.544	1.769	1.00	20.00
	ATOM	13	OG1	THR	2	9.870	67.071	0.713	1.00	20.00
15	ATOM	14	CG2	THR	2	11.672	65.512	1.214	1.00	20.00
	ATOM	15	N	PRO	3	11.385	63.951	3.075	1.00	108.52
	ATOM	16	CA	PRO	3	11.746	62.751	3.877	1.00	109.56
	ATOM	17	C	PRO	3	12.046	62.915	5.329	1.00	110.79
	ATOM	18	O	PRO	3	12.912	63.717	5.675	1.00	114.50
20	ATOM	19	CB	PRO	3	12.862	62.108	3.079	1.00	108.23
	ATOM	20	CG	PRO	3	12.364	62.325	1.691	1.00	109.80
	ATOM	21	CD	PRO	3	11.432	63.520	1.684	1.00	110.35
	ATOM	22	C	CYS	4	12.006	61.576	8.610	1.00	101.54
	ATOM	23	O	CYS	4	10.992	61.129	9.159	1.00	102.94
25	ATOM	24	CB	CYS	4	11.390	63.902	7.916	1.00	96.91
	ATOM	25	SG	CYS	4	12.602	65.259	7.991	1.00	92.39
	ATOM	26	N	CYS	4	11.421	62.190	6.229	1.00	93.44
	ATOM	27	CA	CYS	4	11.991	62.607	7.505	1.00	98.53
	ATOM	28	N	VAL	5	13.247	61.275	8.909	1.00	104.82
30	ATOM	29	CA	VAL	5	13.646	60.294	9.886	1.00	107.58
	ATOM	30	C	VAL	5	13.136	60.620	11.319	1.00	108.57
	ATOM	31	O	VAL	5	13.132	61.786	11.725	1.00	109.82
	ATOM	32	CB	VAL	5	15.156	60.105	9.675	1.00	109.37
	ATOM	33	CG1	VAL	5	15.640	58.854	10.394	1.00	20.00
35	ATOM	34	CG2	VAL	5	15.505	60.050	8.198	1.00	20.00
	ATOM	35	N	PRO	6	12.699	59.575	12.054	1.00	108.52
	ATOM	36	CA	PRO	6	12.163	59.883	13.400	1.00	109.56
	ATOM	37	C	PRO	6	13.031	60.814	14.258	1.00	110.79
	ATOM	38	O	PRO	6	14.244	60.661	14.290	1.00	114.50
40	ATOM	39	CB	PRO	6	11.802	58.510	13.948	1.00	108.23
	ATOM	40	CG	PRO	6	11.267	57.805	12.756	1.00	109.80
	ATOM	41	CD	PRO	6	11.871	58.462	11.534	1.00	110.35
	ATOM	42	N	ALA	7	12.425	61.762	14.969	1.00	109.87
	ATOM	43	CA	ALA	7	13.149	62.757	15.759	1.00	107.78
45	ATOM	44	C	ALA	7	13.359	64.083	15.002	1.00	105.15
	ATOM	45	O	ALA	7	13.160	65.130	15.596	1.00	103.42
	ATOM	46	CB	ALA	7	14.483	62.202	16.214	1.00	109.08
	ATOM	47	N	GLU	8	13.713	64.098	13.677	1.00	101.11
	ATOM	48	CA	GLU	8	14.014	65.418	12.989	1.00	95.31
50	ATOM	49	CB	GLU	8	15.204	65.331	12.069	1.00	92.47
	ATOM	50	CG	GLU	8	15.978	64.029	12.159	1.00	86.82
	ATOM	51	CD	GLU	8	16.286	63.456	10.786	1.00	85.55
	ATOM	52	OE1	GLU	8	15.339	63.266	9.988	1.00	85.72
	ATOM	53	OE2	GLU	8	17.468	63.217	10.502	1.00	81.01
55	ATOM	54	C	GLU	8	12.910	66.092	12.197	1.00	92.66
	ATOM	55	O	GLU	8	11.898	65.476	11.845	1.00	94.43
	ATOM	56	N	CYS	9	13.125	67.372	11.913	1.00	85.19
	ATOM	57	CA	CYS	9	12.146	68.173	11.197	1.00	78.14
	ATOM	58	C	CYS	9	12.824	68.824	10.009	1.00	75.43
60	ATOM	59	O	CYS	9	13.943	69.320	10.114	1.00	76.93
	ATOM	60	CB	CYS	9	11.546	69.243	12.108	1.00	76.59
	ATOM	61	SG	CYS	9	12.706	70.558	12.601	1.00	20.00
	ATOM	62	N	PHE	10	12.113	68.803	8.883	1.00	69.11
	ATOM	63	CA	PHE	10	12.561	69.423	7.612	1.00	62.54
65	ATOM	64	CB	PHE	10	11.556	69.073	6.447	1.00	61.14
	ATOM	65	CG	PHE	10	11.959	69.590	5.084	1.00	61.11

	ATOM	66	CD1	PHE	10	13.042	69.009	4.435	1.00	59.90
	ATOM	67	CD2	PHE	10	11.307	70.663	4.478	1.00	63.23
	ATOM	68	CE1	PHE	10	13.472	69.491	3.198	1.00	61.45
	ATOM	69	CE2	PHE	10	11.725	71.161	3.238	1.00	61.15
5	ATOM	70	CZ	PHE	10	12.808	70.576	2.596	1.00	61.89
	ATOM	71	C	PHE	10	12.671	70.947	7.786	1.00	60.79
	ATOM	72	O	PHE	10	11.680	71.628	8.068	1.00	63.10
	ATOM	73	N	ASP	11	13.884	71.495	7.641	1.00	57.06
	ATOM	74	CA	ASP	11	14.077	72.957	7.749	1.00	52.20
10	ATOM	75	CB	ASP	11	15.493	73.303	8.345	1.00	46.52
	ATOM	76	CG	ASP	11	15.776	74.793	8.611	1.00	47.00
	ATOM	77	OD1	ASP	11	15.241	75.630	7.852	1.00	49.00
	ATOM	78	OD2	ASP	11	16.514	75.095	9.571	1.00	48.43
	ATOM	79	C	ASP	11	13.946	73.545	6.329	1.00	53.02
15	ATOM	80	O	ASP	11	14.740	73.189	5.465	1.00	53.25
	ATOM	81	N	LEU	12	12.968	74.433	6.072	1.00	55.34
	ATOM	82	CA	LEU	12	12.771	75.047	4.739	1.00	54.13
	ATOM	83	C	LEU	12	13.837	76.100	4.385	1.00	54.96
	ATOM	84	O	LEU	12	14.031	76.463	3.228	1.00	54.11
20	ATOM	85	CB	LEU	12	11.345	75.627	4.633	1.00	54.65
	ATOM	86	CG	LEU	12	10.169	74.719	5.032	1.00	20.00
	ATOM	87	CD1	LEU	12	8.864	75.494	4.966	1.00	20.00
	ATOM	88	CD2	LEU	12	10.126	73.495	4.130	1.00	20.00
	ATOM	89	N	LEU	13	14.544	76.591	5.421	1.00	49.64
25	ATOM	90	CA	LEU	13	15.602	77.578	5.241	1.00	46.11
	ATOM	91	CB	LEU	13	15.757	78.492	6.424	1.00	40.39
	ATOM	92	CG	LEU	13	16.931	79.407	6.158	1.00	34.58
	ATOM	93	CD1	LEU	13	16.468	80.667	5.433	1.00	33.04
	ATOM	94	CD2	LEU	13	17.653	79.757	7.443	1.00	33.82
30	ATOM	95	C	LEU	13	16.946	76.917	4.985	1.00	49.82
	ATOM	96	O	LEU	13	17.925	77.617	4.735	1.00	50.94
	ATOM	97	N	VAL	14	17.022	75.584	5.035	1.00	51.20
	ATOM	98	CA	VAL	14	18.284	74.975	4.570	1.00	51.39
	ATOM	99	C	VAL	14	17.956	74.082	3.418	1.00	53.87
35	ATOM	100	O	VAL	14	18.821	73.697	2.629	1.00	50.27
	ATOM	101	CB	VAL	14	19.006	74.025	5.509	1.00	47.82
	ATOM	102	CG1	VAL	14	20.325	73.574	4.888	1.00	20.00
	ATOM	103	CG2	VAL	14	19.242	74.663	6.873	1.00	20.00
	ATOM	104	N	ARG	15	16.665	73.726	3.366	1.00	58.57
40	ATOM	105	CA	ARG	15	16.055	72.900	2.351	1.00	64.33
	ATOM	106	C	ARG	15	16.530	71.444	2.445	1.00	66.69
	ATOM	107	O	ARG	15	16.666	70.723	1.445	1.00	67.11
	ATOM	108	CB	ARG	15	16.270	73.604	0.989	1.00	65.68
	ATOM	109	CG	ARG	15	15.951	75.110	0.964	1.00	69.20
45	ATOM	110	CD	ARG	15	15.056	75.516	-0.203	1.00	20.00
	ATOM	111	NE	ARG	15	14.890	76.979	-0.370	1.00	20.00
	ATOM	112	CZ	ARG	15	14.174	77.536	-1.340	1.00	20.00
	ATOM	113	NH1	ARG	15	13.571	76.772	-2.241	1.00	20.00
	ATOM	114	NH2	ARG	15	14.061	78.856	-1.406	1.00	20.00
50	ATOM	115	N	HIS	16	16.781	71.055	3.710	1.00	69.57
	ATOM	116	CA	HIS	16	17.224	69.720	4.076	1.00	71.62
	ATOM	117	C	HIS	16	16.866	69.319	5.474	1.00	73.99
	ATOM	118	O	HIS	16	16.566	70.153	6.319	1.00	75.45
	ATOM	119	CB	HIS	16	18.725	69.589	4.110	1.00	69.66
55	ATOM	120	CG	HIS	16	19.405	70.049	2.866	1.00	20.00
	ATOM	121	ND1	HIS	16	19.523	69.213	1.775	1.00	20.00
	ATOM	122	CD2	HIS	16	20.001	71.223	2.534	1.00	20.00
	ATOM	123	CE1	HIS	16	20.161	69.856	0.825	1.00	20.00
	ATOM	124	NE2	HIS	16	20.474	71.065	1.255	1.00	20.00
60	ATOM	125	N	CYS	17	16.919	68.029	5.718	1.00	75.75
	ATOM	126	CA	CYS	17	16.557	67.560	7.036	1.00	77.01
	ATOM	127	C	CYS	17	17.602	67.954	8.097	1.00	75.86
	ATOM	128	O	CYS	17	18.796	68.055	7.832	1.00	73.08
	ATOM	129	CB	CYS	17	16.349	66.047	7.008	1.00	80.96
65	ATOM	130	SG	CYS	17	14.681	65.512	7.521	1.00	86.58
	ATOM	131	N	VAL	18	17.094	68.173	9.310	1.00	75.89

	ATOM	132	CA	VAL	18	17.879	68.548	10.452	1.00	77.02
	ATOM	133	C	VAL	18	17.430	67.722	11.616	1.00	78.19
	ATOM	134	O	VAL	18	16.303	67.257	11.683	1.00	77.54
	ATOM	135	CB	VAL	18	17.764	70.048	10.835	1.00	76.68
5	ATOM	136	CG1	VAL	18	17.877	70.230	12.334	1.00	76.28
	ATOM	137	CG2	VAL	18	18.840	70.851	10.134	1.00	76.37
	ATOM	138	N	ALA	19	18.354	67.575	12.568	1.00	80.09
	ATOM	139	CA	ALA	19	18.133	66.822	13.847	1.00	82.16
	ATOM	140	C	ALA	19	17.569	67.717	14.914	1.00	84.28
10	ATOM	141	O	ALA	19	18.140	68.788	15.128	1.00	84.12
	ATOM	142	CB	ALA	19	19.443	66.192	14.323	1.00	80.00
	ATOM	143	N	CYS	20	16.485	67.422	15.578	1.00	87.37
	ATOM	144	CA	CYS	20	16.028	68.542	16.424	1.00	88.33
	ATOM	145	CB	CYS	20	14.594	68.286	16.905	1.00	86.90
15	ATOM	146	SG	CYS	20	13.313	69.181	15.971	1.00	84.56
	ATOM	147	C	CYS	20	16.985	68.911	17.559	1.00	89.64
	ATOM	148	O	CYS	20	16.721	69.839	18.318	1.00	89.77
	ATOM	149	N	GLY	21	18.089	68.176	17.683	1.00	87.37
	ATOM	150	CA	GLY	21	19.018	68.480	18.747	1.00	88.33
20	ATOM	151	C	GLY	21	19.646	69.860	18.549	1.00	89.64
	ATOM	152	O	GLY	21	20.283	70.428	19.433	1.00	89.77
	ATOM	153	N	LEU	22	19.477	70.412	17.342	1.00	87.56
	ATOM	154	CA	LEU	22	20.042	71.712	16.977	1.00	85.33
	ATOM	155	CB	LEU	22	20.266	71.805	15.479	1.00	79.98
25	ATOM	156	CG	LEU	22	21.718	71.871	15.031	1.00	77.91
	ATOM	157	CD1	LEU	22	21.844	71.575	13.549	1.00	78.13
	ATOM	158	CD2	LEU	22	22.318	73.238	15.363	1.00	72.50
	ATOM	159	C	LEU	22	19.173	72.876	17.420	1.00	86.05
	ATOM	160	O	LEU	22	19.687	73.815	18.013	1.00	88.06
30	ATOM	161	N	LEU	23	17.850	72.825	17.121	1.00	85.11
	ATOM	162	CA	LEU	23	16.988	73.968	17.441	1.00	83.33
	ATOM	163	C	LEU	23	16.458	74.012	18.872	1.00	84.21
	ATOM	164	O	LEU	23	15.743	74.963	19.225	1.00	86.28
	ATOM	165	CB	LEU	23	15.799	74.025	16.477	1.00	78.93
35	ATOM	166	CG	LEU	23	15.986	73.794	14.985	1.00	73.68
	ATOM	167	CD1	LEU	23	14.653	73.462	14.352	1.00	20.00
	ATOM	168	CD2	LEU	23	16.630	74.984	14.316	1.00	20.00
	ATOM	169	N	ARG	24	16.794	73.045	19.712	1.00	86.09
	ATOM	170	CA	ARG	24	16.216	73.118	21.057	1.00	85.39
40	ATOM	171	C	ARG	24	17.111	73.811	22.057	1.00	84.10
	ATOM	172	O	ARG	24	16.619	74.341	23.052	1.00	83.24
	ATOM	173	CB	ARG	24	15.804	71.764	21.559	1.00	80.05
	ATOM	174	CG	ARG	24	16.614	70.603	21.059	1.00	20.00
	ATOM	175	CD	ARG	24	16.404	69.446	21.996	1.00	20.00
45	ATOM	176	NE	ARG	24	15.548	68.447	21.411	1.00	20.00
	ATOM	177	CZ	ARG	24	15.437	67.260	21.984	1.00	20.00
	ATOM	178	NH1	ARG	24	16.098	66.996	23.103	1.00	20.00
	ATOM	179	NH2	ARG	24	14.666	66.341	21.439	1.00	20.00
	ATOM	180	N	THR	25	18.441	73.815	21.838	1.00	84.61
50	ATOM	181	CA	THR	25	19.357	74.492	22.785	1.00	85.07
	ATOM	182	C	THR	25	19.272	75.999	22.539	1.00	85.65
	ATOM	183	O	THR	25	19.438	76.822	23.432	1.00	86.77
	ATOM	184	CB	THR	25	20.777	73.909	22.658	1.00	20.00
	ATOM	185	OG1	THR	25	20.698	72.486	22.696	1.00	20.00
55	ATOM	186	CG2	THR	25	21.658	74.413	23.789	1.00	20.00
	ATOM	187	N	PRO	26	19.003	76.308	21.229	1.00	84.19
	ATOM	188	CA	PRO	26	18.604	77.691	20.735	1.00	82.44
	ATOM	189	C	PRO	26	17.178	78.128	21.117	1.00	84.32
	ATOM	190	O	PRO	26	16.770	79.283	20.978	1.00	81.24
60	ATOM	191	CB	PRO	26	18.783	77.600	19.237	1.00	78.00
	ATOM	192	CG	PRO	26	20.010	76.763	19.108	1.00	20.00
	ATOM	193	CD	PRO	26	20.154	75.981	20.392	1.00	20.00
	ATOM	194	N	ARG	27	16.410	77.122	21.601	1.00	87.06
	ATOM	195	CA	ARG	27	14.959	77.115	21.935	1.00	88.17
65	ATOM	196	C	ARG	27	14.204	77.742	20.729	1.00	88.45
	ATOM	197	O	ARG	27	13.116	78.301	20.863	1.00	84.89

	ATOM	198	CB	ARG	27	14.694	77.553	23.416	0.00	86.84
	ATOM	199	CG	ARG	27	15.033	76.419	24.465	0.00	85.55
	ATOM	200	CD	ARG	27	13.977	76.029	25.572	1.00	20.00
	ATOM	201	NE	ARG	27	14.395	74.944	26.507	1.00	20.00
5	ATOM	202	CZ	ARG	27	13.697	74.402	27.516	1.00	20.00
	ATOM	203	NH1	ARG	27	12.477	74.835	27.762	1.00	20.00
	ATOM	204	NH2	ARG	27	14.234	73.451	28.264	1.00	20.00
	ATOM	205	N	PRO	28	14.873	77.581	19.570	1.00	89.49
	ATOM	206	CA	PRO	28	14.220	77.902	18.254	1.00	91.86
10	ATOM	207	C	PRO	28	12.882	77.097	18.097	1.00	95.75
	ATOM	208	O	PRO	28	12.080	77.316	17.192	1.00	96.59
	ATOM	209	CB	PRO	28	15.354	77.698	17.243	0.00	91.23
	ATOM	210	CG	PRO	28	16.536	78.235	18.015	1.00	20.00
	ATOM	211	CD	PRO	28	16.238	78.117	19.483	1.00	20.00
15	ATOM	212	N	LYS	29	12.698	76.148	19.046	1.00	98.41
	ATOM	213	CA	LYS	29	11.582	75.193	19.224	1.00	96.39
	ATOM	214	C	LYS	29	10.652	74.802	18.082	1.00	96.88
	ATOM	215	O	LYS	29	9.804	75.588	17.665	1.00	94.52
	ATOM	216	CB	LYS	29	10.729	75.677	20.396	0.00	95.29
20	ATOM	217	CG	LYS	29	10.916	77.142	20.727	0.00	92.34
	ATOM	218	CD	LYS	29	9.766	77.975	20.196	0.00	89.50
	ATOM	219	CE	LYS	29	9.608	79.268	20.985	1.00	20.00
	ATOM	220	NZ	LYS	29	8.189	79.530	21.345	1.00	20.00
	ATOM	221	N	PRO	30	10.858	73.550	17.537	1.00	97.72
25	ATOM	222	CA	PRO	30	9.868	72.945	16.581	1.00	98.94
	ATOM	223	C	PRO	30	8.507	72.588	17.189	1.00	100.71
	ATOM	224	O	PRO	30	8.446	71.807	18.148	1.00	100.56
	ATOM	225	CB	PRO	30	10.589	71.761	15.992	0.00	97.33
	ATOM	226	CG	PRO	30	11.978	72.258	15.869	1.00	20.00
30	ATOM	227	CD	PRO	30	12.138	73.405	16.838	1.00	20.00
	ATOM	228	N	ALA	31	7.417	73.135	16.646	1.00	101.17
	ATOM	229	CA	ALA	31	6.101	72.844	17.165	1.00	99.22
	ATOM	230	C	ALA	31	5.335	71.906	16.233	1.00	99.01
	ATOM	231	O	ALA	31	4.335	71.296	16.617	1.00	102.17
35	ATOM	232	CB	ALA	31	5.335	74.142	17.377	1.00	20.00
	ATOM	233	N	GLY	32	5.862	71.794	15.028	1.00	95.52
	ATOM	234	CA	GLY	32	5.325	70.856	14.106	1.00	91.44
	ATOM	235	C	GLY	32	6.204	69.621	14.203	1.00	88.49
	ATOM	236	O	GLY	32	7.039	69.386	13.342	1.00	86.70
40	ATOM	237	N	ALA	33	6.002	68.831	15.246	1.00	15.00
	ATOM	238	CA	ALA	33	6.753	67.601	15.416	1.00	15.00
	ATOM	239	C	ALA	33	7.886	67.660	16.411	1.00	15.00
	ATOM	240	O	ALA	33	8.218	66.660	17.052	1.00	15.00
	ATOM	241	CB	ALA	33	7.299	67.152	14.061	1.00	15.00
45	ATOM	242	N	SER	34	8.485	68.810	16.569	1.00	15.00
	ATOM	243	CA	SER	34	9.658	68.894	17.441	1.00	15.00
	ATOM	244	C	SER	34	9.350	68.958	18.934	1.00	15.00
	ATOM	245	O	SER	34	10.099	68.436	19.762	1.00	15.00
	ATOM	246	CB	SER	34	10.483	70.108	17.042	1.00	15.00
50	ATOM	247	OG	SER	34	11.815	69.984	17.514	1.00	20.00
	ATOM	248	N	SER	35	8.253	69.597	19.295	1.00	15.00
	ATOM	249	CA	SER	35	7.875	69.759	20.705	1.00	15.00
	ATOM	250	C	SER	35	8.068	68.437	21.448	1.00	15.00
	ATOM	251	O	SER	35	8.518	68.372	22.592	1.00	15.00
55	ATOM	252	CB	SER	35	6.423	70.249	20.836	1.00	15.00
	ATOM	253	OG	SER	35	5.898	70.621	19.569	1.00	20.00
	ATOM	254	N	PRO	36	7.695	67.392	20.717	1.00	15.00
	ATOM	255	CA	PRO	36	7.832	65.989	21.195	1.00	15.00
	ATOM	256	C	PRO	36	9.196	65.675	21.749	1.00	15.00
60	ATOM	257	O	PRO	36	9.336	65.211	22.881	1.00	15.00
	ATOM	258	CB	PRO	36	7.277	65.194	20.013	1.00	15.00
	ATOM	259	CG	PRO	36	6.151	66.078	19.537	1.00	15.00
	ATOM	260	CD	PRO	36	6.465	67.482	19.918	1.00	15.00
	ATOM	261	N	ALA	37	10.202	65.951	20.940	1.00	15.00
65	ATOM	262	CA	ALA	37	11.583	65.771	21.333	1.00	15.00
	ATOM	263	C	ALA	37	11.893	66.675	22.515	1.00	15.00

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	ATOM	264	O	ALA	37	12.653	66.311	23.403	1.00	15.00
	ATOM	265	CB	ALA	37	12.505	66.075	20.169	1.00	15.00
	ATOM	266	N	PRO	38	11.317	67.902	22.493	1.00	15.00
	ATOM	267	CA	PRO	38	11.559	68.882	23.601	1.00	15.00
5	ATOM	268	C	PRO	38	11.344	68.377	24.975	1.00	15.00
	ATOM	269	O	PRO	38	12.192	68.470	25.851	1.00	15.00
	ATOM	270	CB	PRO	38	10.720	70.069	23.245	1.00	15.00
	ATOM	271	CG	PRO	38	10.931	70.111	21.759	1.00	20.00
	ATOM	272	CD	PRO	38	11.264	68.716	21.280	1.00	20.00
10	ATOM	273	N	ARG	39	10.158	67.808	25.141	1.00	15.00
	ATOM	274	CA	ARG	39	9.759	67.264	26.438	1.00	15.00
	ATOM	275	C	ARG	39	10.345	65.854	26.667	1.00	15.00
	ATOM	276	O	ARG	39	10.096	65.261	27.720	1.00	15.00
	ATOM	277	CB	ARG	39	8.254	67.215	26.577	1.00	15.00
15	ATOM	278	CG	ARG	39	7.561	68.516	26.283	1.00	15.00
	ATOM	279	CD	ARG	39	6.287	68.225	25.523	1.00	15.00
	ATOM	280	NE	ARG	39	5.541	69.430	25.234	1.00	20.00
	ATOM	281	CZ	ARG	39	4.214	69.472	25.163	1.00	20.00
	ATOM	282	NH1	ARG	39	3.504	68.375	25.388	1.00	20.00
20	ATOM	283	NH2	ARG	39	3.602	70.610	24.871	1.00	20.00
	ATOM	284	N	THR	40	11.075	65.325	25.682	1.00	15.00
	ATOM	285	CA	THR	40	11.703	63.978	25.691	1.00	15.00
	ATOM	286	C	THR	40	13.080	64.064	26.358	1.00	15.00
	ATOM	287	O	THR	40	13.453	63.306	27.247	1.00	15.00
25	ATOM	288	CB	THR	40	11.893	63.451	24.288	1.00	15.00
	ATOM	289	OG1	THR	40	10.678	63.674	23.554	1.00	20.00
	ATOM	290	CG2	THR	40	12.213	61.969	24.312	1.00	20.00
	ATOM	291	N	ALA	41	13.813	65.058	25.866	1.00	15.00
	ATOM	292	CA	ALA	41	15.110	65.344	26.414	1.00	15.00
30	ATOM	293	C	ALA	41	14.972	66.152	27.702	1.00	15.00
	ATOM	294	O	ALA	41	15.919	66.244	28.464	1.00	15.00
	ATOM	295	CB	ALA	41	15.976	66.111	25.421	1.00	15.00
	ATOM	296	N	LEU	42	13.832	66.739	27.938	1.00	15.00
	ATOM	297	CA	LEU	42	13.598	67.511	29.152	1.00	15.00
35	ATOM	298	C	LEU	42	13.304	66.561	30.303	1.00	15.00
	ATOM	299	O	LEU	42	13.855	66.696	31.401	1.00	15.00
	ATOM	300	CB	LEU	42	12.434	68.470	28.976	1.00	15.00
	ATOM	301	CG	LEU	42	12.640	69.635	28.006	1.00	20.00
	ATOM	302	CD1	LEU	42	11.399	70.521	27.960	1.00	20.00
40	ATOM	303	CD2	LEU	42	13.862	70.443	28.390	1.00	20.00
	ATOM	304	N	GLN	43	12.442	65.586	30.057	1.00	15.00
	ATOM	305	CA	GLN	43	12.100	64.549	31.044	1.00	15.00
	ATOM	306	C	GLN	43	13.344	63.811	31.479	1.00	15.00
	ATOM	307	O	GLN	43	13.730	63.843	32.645	1.00	15.00
45	ATOM	308	CB	GLN	43	11.093	63.535	30.465	1.00	15.00
	ATOM	309	CG	GLN	43	9.901	64.112	29.739	1.00	20.00
	ATOM	310	CD	GLN	43	9.068	62.991	29.150	1.00	20.00
	ATOM	311	OE1	GLN	43	9.438	61.821	29.197	1.00	20.00
	ATOM	312	NE2	GLN	43	7.886	63.137	28.559	1.00	20.00
50	END					-23.152	-5.163	-22.411	0.00	0.00

TABLE 29

	ATOM	2	CA	PRO	1	-46.326	63.487	-28.122	1.00	98.53
	ATOM	3	C	PRO	1	-45.801	63.556	-26.740	1.00	101.54
5	ATOM	4	O	PRO	1	-44.759	62.925	-26.511	1.00	102.94
	ATOM	5	CB	PRO	1	-45.737	62.321	-28.892	1.00	96.91
	ATOM	6	CG	PRO	1	-46.078	62.724	-30.281	1.00	20.00
	ATOM	7	CD	PRO	1	-46.326	64.223	-30.314	1.00	20.00
	ATOM	8	N	THR	2	-46.338	64.231	-25.789	1.00	104.82
10	ATOM	9	CA	THR	2	-45.584	63.987	-24.602	1.00	107.58
	ATOM	10	C	THR	2	-46.417	63.064	-23.729	1.00	108.57
	ATOM	11	O	THR	2	-47.214	62.308	-24.285	1.00	109.82
	ATOM	12	CB	THR	2	-45.129	65.285	-23.958	1.00	20.00
	ATOM	13	OG1	THR	2	-43.872	65.695	-24.540	1.00	20.00
15	ATOM	14	CG2	THR	2	-44.982	65.100	-22.457	1.00	20.00
	ATOM	15	N	PRO	3	-46.302	63.043	-22.380	1.00	108.52
	ATOM	16	CA	PRO	3	-47.038	61.955	-21.680	1.00	109.56
	ATOM	17	C	PRO	3	-48.469	61.713	-22.027	1.00	110.79
	ATOM	18	O	PRO	3	-49.270	62.645	-21.954	1.00	114.50
20	ATOM	19	CB	PRO	3	-46.792	62.241	-20.213	1.00	108.23
	ATOM	20	CG	PRO	3	-45.367	62.676	-20.256	1.00	109.80
	ATOM	21	CD	PRO	3	-45.057	63.209	-21.641	1.00	110.35
	ATOM	22	C	CYS	4	-51.173	59.446	-22.358	1.00	101.54
	ATOM	23	O	CYS	4	-51.127	58.415	-23.039	1.00	102.94
25	ATOM	24	CB	CYS	4	-50.537	61.286	-23.934	1.00	96.91
	ATOM	25	SG	CYS	4	-51.325	62.925	-23.834	1.00	92.39
	ATOM	26	N	CYS	4	-48.883	60.514	-22.370	1.00	93.44
	ATOM	27	CA	CYS	4	-50.314	60.665	-22.603	1.00	98.53
	ATOM	28	N	VAL	5	-51.984	59.678	-21.354	1.00	104.82
30	ATOM	29	CA	VAL	5	-52.917	58.724	-20.810	1.00	107.58
	ATOM	30	C	VAL	5	-53.968	58.240	-21.849	1.00	108.57
	ATOM	31	O	VAL	5	-54.453	59.037	-22.658	1.00	109.82
	ATOM	32	CB	VAL	5	-53.420	59.353	-19.502	1.00	109.37
	ATOM	33	CG1	VAL	5	-54.137	58.313	-18.653	1.00	20.00
35	ATOM	34	CG2	VAL	5	-52.285	59.997	-18.725	1.00	20.00
	ATOM	35	N	PRO	6	-54.290	56.929	-21.814	1.00	108.52
	ATOM	36	CA	PRO	6	-55.250	56.449	-22.833	1.00	109.56
	ATOM	37	C	PRO	6	-56.514	57.304	-23.002	1.00	110.79
	ATOM	38	O	PRO	6	-57.095	57.738	-22.016	1.00	114.50
40	ATOM	39	CB	PRO	6	-55.406	54.974	-22.491	1.00	108.23
	ATOM	40	CG	PRO	6	-54.032	54.577	-22.094	1.00	109.80
	ATOM	41	CD	PRO	6	-53.320	55.825	-21.621	1.00	110.35
	ATOM	42	N	ALA	7	-56.958	57.535	-24.235	1.00	109.87
	ATOM	43	CA	ALA	7	-58.102	58.398	-24.527	1.00	107.78
45	ATOM	44	C	ALA	7	-57.687	59.839	-24.886	1.00	105.15
	ATOM	45	O	ALA	7	-58.230	60.381	-25.834	1.00	103.42
	ATOM	46	CB	ALA	7	-59.064	58.413	-23.357	1.00	109.08
	ATOM	47	N	GLU	8	-56.694	60.486	-24.196	1.00	101.11
	ATOM	48	CA	GLU	8	-56.382	61.940	-24.507	1.00	95.31
50	ATOM	49	CB	GLU	8	-56.125	62.753	-23.265	1.00	92.47
	ATOM	50	CG	GLU	8	-56.422	62.030	-21.964	1.00	86.82
	ATOM	51	CD	GLU	8	-55.300	62.199	-20.954	1.00	85.55
	ATOM	52	OE1	GLU	8	-54.135	61.885	-21.291	1.00	85.72
	ATOM	53	OE2	GLU	8	-55.580	62.660	-19.838	1.00	81.01
55	ATOM	54	C	GLU	8	-55.246	62.249	-25.462	1.00	92.66
	ATOM	55	O	GLU	8	-54.392	61.402	-25.749	1.00	94.43
	ATOM	56	N	CYS	9	-55.241	63.486	-25.947	1.00	85.19
	ATOM	57	CA	CYS	9	-54.244	63.930	-26.907	1.00	78.14
	ATOM	58	C	CYS	9	-53.596	65.197	-26.386	1.00	75.43
60	ATOM	59	O	CYS	9	-54.270	66.085	-25.870	1.00	76.93
	ATOM	60	CB	CYS	9	-54.880	64.191	-28.271	1.00	76.59
	ATOM	61	SG	CYS	9	-56.004	65.623	-28.324	1.00	20.00
	ATOM	62	N	PHE	10	-52.274	65.246	-26.537	1.00	69.11
	ATOM	63	CA	PHE	10	-51.441	66.408	-26.143	1.00	62.54
65	ATOM	64	CB	PHE	10	-49.909	66.069	-26.312	1.00	61.14
	ATOM	65	CG	PHE	10	-48.964	67.160	-25.862	1.00	61.11

	ATOM	66	CD1	PHE	10	-48.839	67.427	-24.504	1.00	59.90
	ATOM	67	CD2	PHE	10	-48.247	67.938	-26.770	1.00	63.23
	ATOM	68	CE1	PHE	10	-48.012	68.458	-24.055	1.00	61.45
	ATOM	69	CE2	PHE	10	-47.413	68.977	-26.338	1.00	61.15
5	ATOM	70	CZ	PHE	10	-47.295	69.238	-24.981	1.00	61.89
	ATOM	71	C	PHE	10	-51.816	67.632	-26.997	1.00	60.79
	ATOM	72	O	PHE	10	-51.674	67.618	-28.225	1.00	63.10
	ATOM	73	N	ASP	11	-52.321	68.695	-26.358	1.00	57.06
	ATOM	74	CA	ASP	11	-52.670	69.929	-27.093	1.00	52.20
10	ATOM	75	CB	ASP	11	-53.896	70.661	-26.428	1.00	46.52
	ATOM	76	CG	ASP	11	-54.429	71.904	-27.163	1.00	47.00
	ATOM	77	OD1	ASP	11	-53.607	72.601	-27.798	1.00	49.00
	ATOM	78	OD2	ASP	11	-55.650	72.154	-27.099	1.00	48.43
	ATOM	79	C	ASP	11	-51.431	70.848	-27.065	1.00	53.02
15	ATOM	80	O	ASP	11	-51.007	71.239	-25.982	1.00	53.25
	ATOM	81	N	LEU	12	-50.846	71.199	-28.224	1.00	55.34
	ATOM	82	CA	LEU	12	-49.656	72.078	-28.289	1.00	54.13
	ATOM	83	C	LEU	12	-49.965	73.554	-27.984	1.00	54.96
	ATOM	84	O	LEU	12	-49.084	74.350	-27.669	1.00	54.11
20	ATOM	85	CB	LEU	12	-48.958	71.917	-29.656	1.00	54.65
	ATOM	86	CG	LEU	12	-48.652	70.489	-30.143	1.00	20.00
	ATOM	87	CD1	LEU	12	-48.068	70.528	-31.545	1.00	20.00
	ATOM	88	CD2	LEU	12	-47.705	69.799	-29.173	1.00	20.00
	ATOM	89	N	LEU	13	-51.259	73.915	-28.077	1.00	49.64
25	ATOM	90	CA	LEU	13	-51.709	75.272	-27.796	1.00	46.11
	ATOM	91	CB	LEU	13	-52.921	75.665	-28.595	1.00	40.39
	ATOM	92	CG	LEU	13	-53.342	77.050	-28.157	1.00	34.58
	ATOM	93	CD1	LEU	13	-52.632	78.109	-28.993	1.00	33.04
	ATOM	94	CD2	LEU	13	-54.845	77.214	-28.240	1.00	33.82
30	ATOM	95	C	LEU	13	-52.043	75.458	-26.325	1.00	49.82
	ATOM	96	O	LEU	13	-52.362	76.572	-25.915	1.00	50.94
	ATOM	97	N	VAL	14	-51.972	74.398	-25.516	1.00	51.20
	ATOM	98	CA	VAL	14	-52.090	74.662	-24.068	1.00	51.39
	ATOM	99	C	VAL	14	-50.827	74.194	-23.421	1.00	53.87
35	ATOM	100	O	VAL	14	-50.499	74.568	-22.292	1.00	50.27
	ATOM	101	CB	VAL	14	-53.144	73.898	-23.287	1.00	47.82
	ATOM	102	CG1	VAL	14	-53.170	74.371	-21.837	1.00	20.00
	ATOM	103	CG2	VAL	14	-54.521	74.040	-23.926	1.00	20.00
	ATOM	104	N	ARG	15	-50.135	73.320	-24.164	1.00	58.57
40	ATOM	105	CA	ARG	15	-48.867	72.726	-23.811	1.00	64.33
	ATOM	106	C	ARG	15	-49.009	71.738	-22.647	1.00	66.69
	ATOM	107	O	ARG	15	-48.116	71.573	-21.802	1.00	67.11
	ATOM	108	CB	ARG	15	-47.855	73.881	-23.611	1.00	65.68
	ATOM	109	CG	ARG	15	-47.852	74.958	-24.710	1.00	69.20
45	ATOM	110	CD	ARG	15	-46.455	75.282	-25.232	1.00	20.00
	ATOM	111	NE	ARG	15	-46.395	76.448	-26.145	1.00	20.00
	ATOM	112	CZ	ARG	15	-45.271	76.907	-26.685	1.00	20.00
	ATOM	113	NH1	ARG	15	-44.114	76.326	-26.399	1.00	20.00
	ATOM	114	NH2	ARG	15	-45.309	77.945	-27.509	1.00	20.00
50	ATOM	115	N	HIS	16	-50.190	71.091	-22.649	1.00	69.57
	ATOM	116	CA	HIS	16	-50.569	70.089	-21.668	1.00	71.62
	ATOM	117	C	HIS	16	-51.580	69.100	-22.162	1.00	73.99
	ATOM	118	O	HIS	16	-52.273	69.334	-23.144	1.00	75.45
	ATOM	119	CB	HIS	16	-51.290	70.673	-20.480	1.00	69.66
55	ATOM	120	CG	HIS	16	-50.572	71.806	-19.829	1.00	20.00
	ATOM	121	ND1	HIS	16	-49.578	71.573	-18.901	1.00	20.00
	ATOM	122	CD2	HIS	16	-50.693	73.152	-19.952	1.00	20.00
	ATOM	123	CE1	HIS	16	-49.118	72.728	-18.484	1.00	20.00
	ATOM	124	NE2	HIS	16	-49.778	73.700	-19.087	1.00	20.00
60	ATOM	125	N	CYS	17	-51.673	67.995	-21.458	1.00	75.75
	ATOM	126	CA	CYS	17	-52.605	66.977	-21.889	1.00	77.01
	ATOM	127	C	CYS	17	-54.069	67.409	-21.683	1.00	75.86
	ATOM	128	O	CYS	17	-54.410	68.144	-20.761	1.00	73.08
	ATOM	129	CB	CYS	17	-52.313	65.666	-21.164	1.00	80.96
65	ATOM	130	SG	CYS	17	-51.919	64.270	-22.271	1.00	86.58
	ATOM	131	N	VAL	18	-54.917	66.918	-22.586	1.00	75.89

	ATOM	132	CA	VAL	18	-56.328	67.183	-22.590	1.00	77.02
	ATOM	133	C	VAL	18	-57.043	65.891	-22.831	1.00	78.19
	ATOM	134	O	VAL	18	-56.520	64.964	-23.428	1.00	77.54
	ATOM	135	CB	VAL	18	-56.778	68.207	-23.666	1.00	76.68
5	ATOM	136	CG1	VAL	18	-58.164	67.874	-24.178	1.00	76.28
	ATOM	137	CG2	VAL	18	-56.760	69.609	-23.095	1.00	76.37
	ATOM	138	N	ALA	19	-58.295	65.867	-22.366	1.00	80.09
	ATOM	139	CA	ALA	19	-59.226	64.700	-22.516	1.00	82.16
	ATOM	140	C	ALA	19	-59.996	64.781	-23.804	1.00	84.28
10	ATOM	141	O	ALA	19	-60.572	65.839	-24.065	1.00	84.12
	ATOM	142	CB	ALA	19	-60.188	64.635	-21.328	1.00	80.00
	ATOM	143	N	CYS	20	-60.034	63.799	-24.662	1.00	87.37
	ATOM	144	CA	CYS	20	-60.686	64.191	-25.928	1.00	88.33
	ATOM	145	CB	CYS	20	-60.405	63.142	-27.010	1.00	86.90
15	ATOM	146	SG	CYS	20	-59.085	63.597	-28.179	1.00	84.56
	ATOM	147	C	CYS	20	-62.171	64.535	-25.796	1.00	89.64
	ATOM	148	O	CYS	20	-62.816	64.893	-26.777	1.00	89.77
	ATOM	149	N	GLY	21	-62.716	64.413	-24.587	1.00	87.37
	ATOM	150	CA	GLY	21	-64.118	64.716	-24.415	1.00	88.33
20	ATOM	151	C	GLY	21	-64.395	66.197	-24.676	1.00	89.64
	ATOM	152	O	GLY	21	-65.532	66.641	-24.817	1.00	89.77
	ATOM	153	N	LEU	22	-63.321	66.993	-24.726	1.00	87.56
	ATOM	154	CA	LEU	22	-63.413	68.439	-24.935	1.00	85.33
	ATOM	155	CB	LEU	22	-62.217	69.152	-24.329	1.00	79.98
25	ATOM	156	CG	LEU	22	-62.514	70.044	-23.134	1.00	77.91
	ATOM	157	CD1	LEU	22	-61.242	70.390	-22.383	1.00	78.13
	ATOM	158	CD2	LEU	22	-63.240	71.313	-23.583	1.00	72.50
	ATOM	159	C	LEU	22	-63.523	68.817	-26.401	1.00	86.05
	ATOM	160	O	LEU	22	-64.389	69.606	-26.752	1.00	88.06
30	ATOM	161	N	LEU	23	-62.634	68.262	-27.263	1.00	85.11
	ATOM	162	CA	LEU	23	-62.638	68.669	-28.672	1.00	83.33
	ATOM	163	C	LEU	23	-63.646	67.948	-29.564	1.00	84.21
	ATOM	164	O	LEU	23	-63.726	68.258	-30.763	1.00	86.28
	ATOM	165	CB	LEU	23	-61.240	68.501	-29.279	1.00	78.93
35	ATOM	166	CG	LEU	23	-59.996	68.931	-28.517	1.00	73.68
	ATOM	167	CD1	LEU	23	-58.779	68.263	-29.116	1.00	20.00
	ATOM	168	CD2	LEU	23	-59.847	70.433	-28.503	1.00	20.00
	ATOM	169	N	ARG	24	-64.431	67.025	-29.029	1.00	86.09
	ATOM	170	CA	ARG	24	-65.345	66.336	-29.944	1.00	85.39
40	ATOM	171	C	ARG	24	-66.719	66.961	-30.002	1.00	84.10
	ATOM	172	O	ARG	24	-67.419	66.806	-31.002	1.00	83.24
	ATOM	173	CB	ARG	24	-65.439	64.869	-29.636	1.00	80.05
	ATOM	174	CG	ARG	24	-65.252	64.487	-28.197	1.00	20.00
	ATOM	175	CD	ARG	24	-65.843	63.120	-27.993	1.00	20.00
45	ATOM	176	NE	ARG	24	-64.817	62.118	-27.866	1.00	20.00
	ATOM	177	CZ	ARG	24	-65.133	60.903	-27.451	1.00	20.00
	ATOM	178	NH1	ARG	24	-66.394	60.601	-27.171	1.00	20.00
	ATOM	179	NH2	ARG	24	-64.191	59.992	-27.319	1.00	20.00
	ATOM	180	N	THR	25	-67.153	67.665	-28.938	1.00	84.61
50	ATOM	181	CA	THR	25	-68.489	68.305	-28.953	1.00	85.07
	ATOM	182	C	THR	25	-68.403	69.572	-29.805	1.00	85.65
	ATOM	183	O	THR	25	-69.354	69.998	-30.449	1.00	86.77
	ATOM	184	CB	THR	25	-68.979	68.544	-27.512	1.00	20.00
	ATOM	185	OG1	THR	25	-68.815	67.341	-26.763	1.00	20.00
55	ATOM	186	CG2	THR	25	-70.440	68.963	-27.514	1.00	20.00
	ATOM	187	N	PRO	26	-67.163	70.161	-29.756	1.00	84.19
	ATOM	188	CA	PRO	26	-66.699	71.269	-30.690	1.00	82.44
	ATOM	189	C	PRO	26	-66.412	70.819	-32.134	1.00	84.32
	ATOM	190	O	PRO	26	-66.228	71.611	-33.061	1.00	81.24
60	ATOM	191	CB	PRO	26	-65.461	71.810	-30.013	1.00	78.00
	ATOM	192	CG	PRO	26	-65.830	71.754	-28.570	1.00	20.00
	ATOM	193	CD	PRO	26	-66.935	70.733	-28.432	1.00	20.00
	ATOM	194	N	ARG	27	-66.362	69.473	-32.287	1.00	87.06
	ATOM	195	CA	ARG	27	-65.972	68.669	-33.478	1.00	88.17
65	ATOM	196	C	ARG	27	-64.632	69.251	-34.011	1.00	88.45
	ATOM	197	O	ARG	27	-64.301	69.146	-35.190	1.00	84.89

	ATOM	198	CB	ARG	27	-67.193	68.373	-34.414	0.00	86.84
	ATOM	199	CG	ARG	27	-68.144	67.242	-33.850	0.00	85.55
	ATOM	200	CD	ARG	27	-68.573	66.039	-34.778	1.00	20.00
	ATOM	201	NE	ARG	27	-69.466	65.025	-34.146	1.00	20.00
5	ATOM	202	CZ	ARG	27	-69.961	63.901	-34.686	1.00	20.00
	ATOM	203	NH1	ARG	27	-69.652	63.594	-35.929	1.00	20.00
	ATOM	204	NH2	ARG	27	-70.762	63.118	-33.980	1.00	20.00
	ATOM	205	N	PRO	28	-63.913	69.846	-33.037	1.00	89.49
	ATOM	206	CA	PRO	28	-62.490	70.267	-33.280	1.00	91.86
10	ATOM	207	C	PRO	28	-61.633	69.045	-33.765	1.00	95.75
	ATOM	208	O	PRO	28	-60.489	69.168	-34.195	1.00	96.59
	ATOM	209	CB	PRO	28	-62.115	70.992	-31.982	0.00	91.23
	ATOM	210	CG	PRO	28	-63.406	71.706	-31.660	1.00	20.00
	ATOM	211	CD	PRO	28	-64.538	70.950	-32.296	1.00	20.00
15	ATOM	212	N	LYS	29	-62.271	67.853	-33.666	1.00	98.41
	ATOM	213	CA	LYS	29	-61.796	66.495	-34.008	1.00	96.39
	ATOM	214	C	LYS	29	-60.315	66.149	-34.105	1.00	96.88
	ATOM	215	O	LYS	29	-59.639	66.536	-35.056	1.00	94.52
	ATOM	216	CB	LYS	29	-62.475	66.071	-35.309	0.00	95.29
20	ATOM	217	CG	LYS	29	-63.017	67.227	-36.124	0.00	92.34
	ATOM	218	CD	LYS	29	-62.106	67.550	-37.290	0.00	89.50
	ATOM	219	CE	LYS	29	-62.868	68.243	-38.413	1.00	20.00
	ATOM	220	NZ	LYS	29	-62.546	67.662	-39.744	1.00	20.00
	ATOM	221	N	PRO	30	-59.794	65.426	-33.049	1.00	97.72
25	ATOM	222	CA	PRO	30	-58.424	64.812	-33.128	1.00	98.94
	ATOM	223	C	PRO	30	-58.276	63.670	-34.140	1.00	100.71
	ATOM	224	O	PRO	30	-59.000	62.669	-34.049	1.00	100.56
	ATOM	225	CB	PRO	30	-58.114	64.401	-31.713	0.00	97.33
	ATOM	226	CG	PRO	30	-58.715	65.497	-30.920	1.00	20.00
30	ATOM	227	CD	PRO	30	-59.768	66.156	-31.779	1.00	20.00
	ATOM	228	N	ALA	31	-57.350	63.795	-35.093	1.00	101.17
	ATOM	229	CA	ALA	31	-57.153	62.759	-36.080	1.00	99.22
	ATOM	230	C	ALA	31	-55.873	61.972	-35.806	1.00	99.01
	ATOM	231	O	ALA	31	-55.670	60.874	-36.328	1.00	102.17
35	ATOM	232	CB	ALA	31	-57.125	63.375	-37.471	1.00	20.00
	ATOM	233	N	GLY	32	-55.052	62.557	-34.953	1.00	95.52
	ATOM	234	CA	GLY	32	-53.888	61.874	-34.512	1.00	91.44
	ATOM	235	C	GLY	32	-54.247	61.252	-33.173	1.00	88.49
	ATOM	236	O	GLY	32	-53.859	61.758	-32.131	1.00	86.70
40	ATOM	237	N	ALA	33	-54.977	60.147	-33.210	1.00	15.00
	ATOM	238	CA	ALA	33	-55.340	59.443	-31.994	1.00	15.00
	ATOM	239	C	ALA	33	-56.751	59.668	-31.507	1.00	15.00
	ATOM	240	O	ALA	33	-57.355	58.787	-30.891	1.00	15.00
	ATOM	241	CB	ALA	33	-54.360	59.818	-30.883	1.00	15.00
45	ATOM	242	N	SER	34	-57.299	60.823	-31.774	1.00	15.00
	ATOM	243	CA	SER	34	-58.623	61.131	-31.230	1.00	15.00
	ATOM	244	C	SER	34	-59.793	60.508	-31.986	1.00	15.00
	ATOM	245	O	SER	34	-60.812	60.142	-31.396	1.00	15.00
	ATOM	246	CB	SER	34	-58.798	62.641	-31.182	1.00	15.00
50	ATOM	247	OG	SER	34	-59.823	62.997	-30.269	1.00	20.00
	ATOM	248	N	SER	35	-59.666	60.383	-33.294	1.00	15.00
	ATOM	249	CA	SER	35	-60.740	59.837	-34.132	1.00	15.00
	ATOM	250	C	SER	35	-61.333	58.593	-33.469	1.00	15.00
	ATOM	251	O	SER	35	-62.539	58.346	-33.469	1.00	15.00
55	ATOM	252	CB	SER	35	-60.228	59.507	-35.544	1.00	15.00
	ATOM	253	OG	SER	35	-58.914	60.012	-35.737	1.00	20.00
	ATOM	254	N	PRO	36	-60.400	57.833	-32.905	1.00	15.00
	ATOM	255	CA	PRO	36	-60.726	56.593	-32.150	1.00	15.00
	ATOM	256	C	PRO	36	-61.816	56.781	-31.130	1.00	15.00
60	ATOM	257	O	PRO	36	-62.821	56.069	-31.128	1.00	15.00
	ATOM	258	CB	PRO	36	-59.341	56.109	-31.720	1.00	15.00
	ATOM	259	CG	PRO	36	-58.495	56.466	-32.916	1.00	15.00
	ATOM	260	CD	PRO	36	-59.133	57.614	-33.616	1.00	15.00
	ATOM	261	N	ALA	37	-61.612	57.762	-30.271	1.00	15.00
65	ATOM	262	CA	ALA	37	-62.584	58.124	-29.263	1.00	15.00
	ATOM	263	C	ALA	37	-63.867	58.581	-29.939	1.00	15.00

	ATOM	264	O	ALA	37	-64.960	58.327	-29.452	1.00	15.00
	ATOM	265	CB	ALA	37	-62.033	59.216	-28.367	1.00	15.00
	ATOM	266	N	PRO	38	-63.715	59.312	-31.071	1.00	15.00
	ATOM	267	CA	PRO	38	-64.908	59.825	-31.819	1.00	15.00
5	ATOM	268	C	PRO	38	-65.954	58.827	-32.138	1.00	15.00
	ATOM	269	O	PRO	38	-67.130	58.989	-31.844	1.00	15.00
	ATOM	270	CB	PRO	38	-64.336	60.519	-33.015	1.00	15.00
	ATOM	271	CG	PRO	38	-63.139	61.178	-32.394	1.00	20.00
	ATOM	272	CD	PRO	38	-62.719	60.376	-31.183	1.00	20.00
10	ATOM	273	N	ARG	39	-65.478	57.752	-32.751	1.00	15.00
	ATOM	274	CA	ARG	39	-66.365	56.665	-33.161	1.00	15.00
	ATOM	275	C	ARG	39	-66.683	55.717	-31.984	1.00	15.00
	ATOM	276	O	ARG	39	-67.421	54.747	-32.174	1.00	15.00
	ATOM	277	CB	ARG	39	-65.774	55.871	-34.305	1.00	15.00
15	ATOM	278	CG	ARG	39	-65.337	56.704	-35.478	1.00	15.00
	ATOM	279	CD	ARG	39	-64.041	56.141	-36.013	1.00	15.00
	ATOM	280	NE	ARG	39	-63.572	56.869	-37.173	1.00	20.00
	ATOM	281	CZ	ARG	39	-62.892	56.307	-38.167	1.00	20.00
	ATOM	282	NH1	ARG	39	-62.631	55.007	-38.138	1.00	20.00
20	ATOM	283	NH2	ARG	39	-62.475	57.045	-39.185	1.00	20.00
	ATOM	284	N	THR	40	-66.104	55.981	-30.810	1.00	15.00
	ATOM	285	CA	THR	40	-66.256	55.182	-29.566	1.00	15.00
	ATOM	286	C	THR	40	-67.496	55.659	-28.804	1.00	15.00
	ATOM	287	O	THR	40	-68.365	54.906	-28.378	1.00	15.00
25	ATOM	288	CB	THR	40	-65.058	55.342	-28.660	1.00	15.00
	ATOM	289	OG1	THR	40	-63.869	55.215	-29.458	1.00	20.00
	ATOM	290	CG2	THR	40	-65.062	54.286	-27.573	1.00	20.00
	ATOM	291	N	ALA	41	-67.522	56.982	-28.668	1.00	15.00
	ATOM	292	CA	ALA	41	-68.643	57.626	-28.042	1.00	15.00
30	ATOM	293	C	ALA	41	-69.797	57.758	-29.033	1.00	15.00
	ATOM	294	O	ALA	41	-70.919	58.005	-28.627	1.00	15.00
	ATOM	295	CB	ALA	41	-68.267	59.005	-27.511	1.00	15.00
	ATOM	296	N	LEU	42	-69.534	57.615	-30.302	1.00	15.00
	ATOM	297	CA	LEU	42	-70.573	57.699	-31.320	1.00	15.00
35	ATOM	298	C	LEU	42	-71.336	56.384	-31.369	1.00	15.00
	ATOM	299	O	LEU	42	-72.571	56.361	-31.394	1.00	15.00
	ATOM	300	CB	LEU	42	-69.980	57.993	-32.687	1.00	15.00
	ATOM	301	CG	LEU	42	-69.358	59.376	-32.887	1.00	20.00
	ATOM	302	CD1	LEU	42	-68.834	59.528	-34.311	1.00	20.00
40	ATOM	303	CD2	LEU	42	-70.359	60.465	-32.564	1.00	20.00
	ATOM	304	N	GLN	43	-70.606	55.279	-31.371	1.00	15.00
	ATOM	305	CA	GLN	43	-71.193	53.930	-31.352	1.00	15.00
	ATOM	306	C	GLN	43	-72.076	53.761	-30.138	1.00	15.00
	ATOM	307	O	GLN	43	-73.281	53.553	-30.249	1.00	15.00
45	ATOM	308	CB	GLN	43	-70.099	52.843	-31.331	1.00	15.00
	ATOM	309	CG	GLN	43	-68.968	53.009	-32.318	1.00	20.00
	ATOM	310	CD	GLN	43	-67.935	51.921	-32.101	1.00	20.00
	ATOM	311	OE1	GLN	43	-68.019	51.131	-31.165	1.00	20.00
	ATOM	312	NE2	GLN	43	-66.879	51.695	-32.877	1.00	20.00
50	END					0.000	0.000	0.000	0.00	0.00

TABLE 30

	ATOM	2	CA	PRO	1	-62.683	100.238	-0.912	1.00	98.53
	ATOM	3	C	PRO	1	-63.677	101.230	-1.379	1.00	101.54
5	ATOM	4	O	PRO	1	-63.750	102.287	-0.736	1.00	102.94
	ATOM	5	CB	PRO	1	-62.238	100.512	0.511	1.00	96.91
	ATOM	6	CG	PRO	1	-61.003	99.687	0.565	1.00	20.00
	ATOM	7	CD	PRO	1	-60.514	99.437	-0.853	1.00	20.00
	ATOM	8	N	THR	2	-64.440	101.073	-2.400	1.00	104.82
10	ATOM	9	CA	THR	2	-65.300	102.208	-2.480	1.00	107.58
	ATOM	10	C	THR	2	-66.677	101.754	-2.024	1.00	108.57
	ATOM	11	O	THR	2	-66.741	100.813	-1.233	1.00	109.82
	ATOM	12	CB	THR	2	-65.232	102.860	-3.849	1.00	20.00
	ATOM	13	OG1	THR	2	-64.150	103.818	-3.873	1.00	20.00
15	ATOM	14	CG2	THR	2	-66.551	103.544	-4.169	1.00	20.00
	ATOM	15	N	PRO	3	-67.815	102.353	-2.447	1.00	108.52
	ATOM	16	CA	PRO	3	-69.068	101.926	-1.767	1.00	109.56
	ATOM	17	C	PRO	3	-69.343	100.469	-1.607	1.00	110.79
	ATOM	18	O	PRO	3	-69.340	99.745	-2.602	1.00	114.50
20	ATOM	19	CB	PRO	3	-70.152	102.688	-2.500	1.00	108.23
	ATOM	20	CG	PRO	3	-69.471	103.998	-2.708	1.00	109.80
	ATOM	21	CD	PRO	3	-67.972	103.780	-2.694	1.00	110.35
	ATOM	22	C	CYS	4	-70.800	97.840	0.270	1.00	101.54
	ATOM	23	O	CYS	4	-70.568	97.640	1.469	1.00	102.94
25	ATOM	24	CB	CYS	4	-68.548	97.850	-0.826	1.00	96.91
	ATOM	25	SG	CYS	4	-68.308	97.145	-2.488	1.00	92.39
	ATOM	26	N	CYS	4	-69.621	99.964	-0.427	1.00	93.44
	ATOM	27	CA	CYS	4	-69.852	98.545	-0.672	1.00	98.53
	ATOM	28	N	VAL	5	-71.863	97.449	-0.391	1.00	104.82
30	ATOM	29	CA	VAL	5	-72.999	96.782	0.193	1.00	107.58
	ATOM	30	C	VAL	5	-72.632	95.427	0.862	1.00	108.57
	ATOM	31	O	VAL	5	-71.806	94.678	0.333	1.00	109.82
	ATOM	32	CB	VAL	5	-74.076	96.787	-0.902	1.00	109.37
	ATOM	33	CG1	VAL	5	-75.434	96.433	-0.314	1.00	20.00
35	ATOM	34	CG2	VAL	5	-74.129	98.124	-1.620	1.00	20.00
	ATOM	35	N	PRO	6	-73.248	95.146	2.030	1.00	108.52
	ATOM	36	CA	PRO	6	-72.866	93.883	2.701	1.00	109.56
	ATOM	37	C	PRO	6	-72.839	92.640	1.799	1.00	110.79
	ATOM	38	O	PRO	6	-73.734	92.456	0.986	1.00	114.50
40	ATOM	39	CB	PRO	6	-73.751	93.869	3.938	1.00	108.23
	ATOM	40	CG	PRO	6	-73.772	95.295	4.350	1.00	109.80
	ATOM	41	CD	PRO	6	-73.486	96.125	3.118	1.00	110.35
	ATOM	42	N	ALA	7	-71.838	91.777	1.947	1.00	109.87
	ATOM	43	CA	ALA	7	-71.660	90.602	1.095	1.00	107.78
45	ATOM	44	C	ALA	7	-70.684	90.851	-0.072	1.00	105.15
	ATOM	45	O	ALA	7	-69.850	89.997	-0.324	1.00	103.42
	ATOM	46	CB	ALA	7	-72.997	90.131	0.560	1.00	109.08
	ATOM	47	N	GLU	8	-70.709	92.023	-0.782	1.00	101.11
	ATOM	48	CA	GLU	8	-69.805	92.195	-1.991	1.00	95.31
50	ATOM	49	CB	GLU	8	-70.499	92.882	-3.138	1.00	92.47
	ATOM	50	CG	GLU	8	-71.990	93.083	-2.945	1.00	86.82
	ATOM	51	CD	GLU	8	-72.423	94.495	-3.301	1.00	85.55
	ATOM	52	OE1	GLU	8	-71.848	95.458	-2.745	1.00	85.72
	ATOM	53	OE2	GLU	8	-73.319	94.638	-4.146	1.00	81.01
55	ATOM	54	C	GLU	8	-68.477	92.904	-1.808	1.00	92.66
	ATOM	55	O	GLU	8	-68.245	93.598	-0.812	1.00	94.43
	ATOM	56	N	CYS	9	-67.606	92.726	-2.796	1.00	85.19
	ATOM	57	CA	CYS	9	-66.272	93.304	-2.755	1.00	78.14
	ATOM	58	C	CYS	9	-66.043	94.091	-4.028	1.00	75.43
60	ATOM	59	O	CYS	9	-66.398	93.647	-5.118	1.00	76.93
	ATOM	60	CB	CYS	9	-65.211	92.213	-2.616	1.00	76.59
	ATOM	61	SG	CYS	9	-65.028	91.141	-4.076	1.00	20.00
	ATOM	62	N	PHE	10	-65.444	95.267	-3.853	1.00	69.11
	ATOM	63	CA	PHE	10	-65.079	96.181	-4.963	1.00	62.54
65	ATOM	64	CB	PHE	10	-64.535	97.548	-4.393	1.00	61.14
	ATOM	65	CG	PHE	10	-64.206	98.587	-5.441	1.00	61.11

	ATOM	66	CD1	PHE	10	-65.244	99.197	-6.136	1.00	59.90
	ATOM	67	CD2	PHE	10	-62.892	98.921	-5.765	1.00	63.23
	ATOM	68	CE1	PHE	10	-64.977	100.126	-7.143	1.00	61.45
	ATOM	69	CE2	PHE	10	-62.605	99.850	-6.773	1.00	61.15
5	ATOM	70	CZ	PHE	10	-63.647	100.453	-7.463	1.00	61.89
	ATOM	71	C	PHE	10	-64.022	95.516	-5.861	1.00	60.79
	ATOM	72	O	PHE	10	-62.914	95.201	-5.413	1.00	63.10
	ATOM	73	N	ASP	11	-64.362	95.273	-7.133	1.00	57.06
	ATOM	74	CA	ASP	11	-63.395	94.675	-8.078	1.00	52.20
10	ATOM	75	CB	ASP	11	-64.123	93.772	-9.144	1.00	46.52
	ATOM	76	CG	ASP	11	-63.216	93.002	-10.121	1.00	47.00
	ATOM	77	OD1	ASP	11	-62.132	93.534	-10.448	1.00	49.00
	ATOM	78	OD2	ASP	11	-63.596	91.887	-10.531	1.00	48.43
	ATOM	79	C	ASP	11	-62.663	95.835	-8.785	1.00	53.02
15	ATOM	80	O	ASP	11	-63.316	96.622	-9.463	1.00	53.25
	ATOM	81	N	LEU	12	-61.331	95.956	-8.643	1.00	55.34
	ATOM	82	CA	LEU	12	-60.550	97.037	-9.287	1.00	54.13
	ATOM	83	C	LEU	12	-60.384	96.854	-10.805	1.00	54.96
	ATOM	84	O	LEU	12	-60.067	97.786	-11.541	1.00	54.11
20	ATOM	85	CB	LEU	12	-59.185	97.191	-8.582	1.00	54.65
	ATOM	86	CG	LEU	12	-59.178	97.304	-7.048	1.00	20.00
	ATOM	87	CD1	LEU	12	-57.749	97.338	-6.532	1.00	20.00
	ATOM	88	CD2	LEU	12	-59.947	98.543	-6.613	1.00	20.00
	ATOM	89	N	LEU	13	-60.612	95.613	-11.276	1.00	49.64
25	ATOM	90	CA	LEU	13	-60.516	95.290	-12.694	1.00	46.11
	ATOM	91	CB	LEU	13	-60.093	93.868	-12.944	1.00	40.39
	ATOM	92	CG	LEU	13	-60.114	93.630	-14.437	1.00	34.58
	ATOM	93	CD1	LEU	13	-58.762	93.982	-15.048	1.00	33.04
	ATOM	94	CD2	LEU	13	-60.494	92.199	-14.755	1.00	33.82
30	ATOM	95	C	LEU	13	-61.839	95.514	-13.410	1.00	49.82
	ATOM	96	O	LEU	13	-61.898	95.361	-14.628	1.00	50.94
	ATOM	97	N	VAL	14	-62.901	95.879	-12.688	1.00	51.20
	ATOM	98	CA	VAL	14	-64.101	96.294	-13.441	1.00	51.39
	ATOM	99	C	VAL	14	-64.402	97.709	-13.063	1.00	53.87
35	ATOM	100	O	VAL	14	-65.133	98.423	-13.754	1.00	50.27
	ATOM	101	CB	VAL	14	-65.415	95.600	-13.131	1.00	47.82
	ATOM	102	CG1	VAL	14	-66.509	96.103	-14.067	1.00	20.00
	ATOM	103	CG2	VAL	14	-65.278	94.085	-13.222	1.00	20.00
	ATOM	104	N	ARG	15	-63.839	98.087	-11.907	1.00	58.57
40	ATOM	105	CA	ARG	15	-63.929	99.399	-11.312	1.00	64.33
	ATOM	106	C	ARG	15	-65.346	99.695	-10.805	1.00	66.69
	ATOM	107	O	ARG	15	-65.835	100.834	-10.824	1.00	67.11
	ATOM	108	CB	ARG	15	-63.338	100.410	-12.326	1.00	65.68
	ATOM	109	CG	ARG	15	-61.993	100.007	-12.955	1.00	69.20
45	ATOM	110	CD	ARG	15	-60.947	101.117	-12.900	1.00	20.00
	ATOM	111	NE	ARG	15	-59.711	100.835	-13.668	1.00	20.00
	ATOM	112	CZ	ARG	15	-58.693	101.683	-13.768	1.00	20.00
	ATOM	113	NH1	ARG	15	-58.758	102.868	-13.176	1.00	20.00
	ATOM	114	NH2	ARG	15	-57.614	101.343	-14.460	1.00	20.00
50	ATOM	115	N	HIS	16	-65.982	98.597	-10.354	1.00	69.57
	ATOM	116	CA	HIS	16	-67.326	98.605	-9.802	1.00	71.62
	ATOM	117	C	HIS	16	-67.601	97.488	-8.842	1.00	73.99
	ATOM	118	O	HIS	16	-66.900	96.485	-8.815	1.00	75.45
	ATOM	119	CB	HIS	16	-68.389	98.364	-10.842	1.00	69.66
55	ATOM	120	CG	HIS	16	-68.298	99.265	-12.027	1.00	20.00
	ATOM	121	ND1	HIS	16	-68.849	100.529	-11.998	1.00	20.00
	ATOM	122	CD2	HIS	16	-67.743	99.101	-13.255	1.00	20.00
	ATOM	123	CE1	HIS	16	-68.634	101.104	-13.157	1.00	20.00
	ATOM	124	NE2	HIS	16	-67.982	100.266	-13.942	1.00	20.00
60	ATOM	125	N	CYS	17	-68.645	97.663	-8.064	1.00	75.75
	ATOM	126	CA	CYS	17	-68.958	96.643	-7.089	1.00	77.01
	ATOM	127	C	CYS	17	-69.479	95.352	-7.749	1.00	75.86
	ATOM	128	O	CYS	17	-70.127	95.367	-8.791	1.00	73.08
	ATOM	129	CB	CYS	17	-69.964	97.185	-6.077	1.00	80.96
65	ATOM	130	SG	CYS	17	-69.376	97.155	-4.349	1.00	86.58
	ATOM	131	N	VAL	18	-69.162	94.236	-7.093	1.00	75.89

	ATOM	132	CA	VAL	18	-69.544	92.919	-7.521	1.00	77.02
	ATOM	133	C	VAL	18	-70.048	92.171	-6.327	1.00	78.19
	ATOM	134	O	VAL	18	-69.689	92.445	-5.193	1.00	77.54
	ATOM	135	CB	VAL	18	-68.391	92.104	-8.166	1.00	76.68
5	ATOM	136	CG1	VAL	18	-68.540	90.628	-7.860	1.00	76.28
	ATOM	137	CG2	VAL	18	-68.371	92.322	-9.664	1.00	76.37
	ATOM	138	N	ALA	19	-70.885	91.174	-6.623	1.00	80.09
	ATOM	139	CA	ALA	19	-71.497	90.258	-5.604	1.00	82.16
	ATOM	140	C	ALA	19	-70.613	89.072	-5.340	1.00	84.28
10	ATOM	141	O	ALA	19	-70.199	88.436	-6.311	1.00	84.12
	ATOM	142	CB	ALA	19	-72.878	89.795	-6.071	1.00	80.00
	ATOM	143	N	CYS	20	-70.239	88.728	-4.139	1.00	87.37
	ATOM	144	CA	CYS	20	-69.221	87.658	-4.157	1.00	88.33
	ATOM	145	CB	CYS	20	-68.567	87.531	-2.776	1.00	86.90
15	ATOM	146	SG	CYS	20	-66.939	88.333	-2.633	1.00	84.56
	ATOM	147	C	CYS	20	-69.716	86.322	-4.713	1.00	89.64
	ATOM	148	O	CYS	20	-68.955	85.362	-4.796	1.00	89.77
	ATOM	149	N	GLY	21	-70.994	86.254	-5.081	1.00	87.37
	ATOM	150	CA	GLY	21	-71.511	85.010	-5.601	1.00	88.33
20	ATOM	151	C	GLY	21	-70.841	84.650	-6.928	1.00	89.64
	ATOM	152	O	GLY	21	-70.944	83.538	-7.440	1.00	89.77
	ATOM	153	N	LEU	22	-70.142	85.629	-7.515	1.00	87.56
	ATOM	154	CA	LEU	22	-69.467	85.461	-8.803	1.00	85.33
	ATOM	155	CB	LEU	22	-69.326	86.792	-9.519	1.00	79.98
25	ATOM	156	CG	LEU	22	-70.139	86.946	-10.795	1.00	77.91
	ATOM	157	CD1	LEU	22	-70.231	88.403	-11.209	1.00	78.13
	ATOM	158	CD2	LEU	22	-69.536	86.101	-11.918	1.00	72.50
	ATOM	159	C	LEU	22	-68.096	84.823	-8.670	1.00	86.05
	ATOM	160	O	LEU	22	-67.799	83.885	-9.398	1.00	88.06
30	ATOM	161	N	LEU	23	-67.247	85.340	-7.746	1.00	85.11
	ATOM	162	CA	LEU	23	-65.878	84.822	-7.647	1.00	83.33
	ATOM	163	C	LEU	23	-65.711	83.562	-6.801	1.00	84.21
	ATOM	164	O	LEU	23	-64.586	83.049	-6.693	1.00	86.28
	ATOM	165	CB	LEU	23	-64.938	85.903	-7.104	1.00	78.93
35	ATOM	166	CG	LEU	23	-65.018	87.337	-7.605	1.00	73.68
	ATOM	167	CD1	LEU	23	-64.326	88.256	-6.622	1.00	20.00
	ATOM	168	CD2	LEU	23	-64.434	87.475	-8.990	1.00	20.00
	ATOM	169	N	ARG	24	-66.777	83.029	-6.224	1.00	86.09
	ATOM	170	CA	ARG	24	-66.545	81.848	-5.388	1.00	85.39
40	ATOM	171	C	ARG	24	-66.736	80.544	-6.127	1.00	84.10
	ATOM	172	O	ARG	24	-66.164	79.529	-5.732	1.00	83.24
	ATOM	173	CB	ARG	24	-67.378	81.879	-4.138	1.00	80.05
	ATOM	174	CG	ARG	24	-68.701	82.579	-4.248	1.00	20.00
	ATOM	175	CD	ARG	24	-69.576	82.109	-3.119	1.00	20.00
45	ATOM	176	NE	ARG	24	-69.700	83.115	-2.098	1.00	20.00
	ATOM	177	CZ	ARG	24	-70.609	82.977	-1.147	1.00	20.00
	ATOM	178	NH1	ARG	24	-71.391	81.906	-1.123	1.00	20.00
	ATOM	179	NH2	ARG	24	-70.734	83.907	-0.222	1.00	20.00
	ATOM	180	N	THR	25	-67.551	80.524	-7.200	1.00	84.61
50	ATOM	181	CA	THR	25	-67.760	79.272	-7.963	1.00	85.07
	ATOM	182	C	THR	25	-66.532	79.036	-8.843	1.00	85.65
	ATOM	183	O	THR	25	-66.143	77.914	-9.143	1.00	86.77
	ATOM	184	CB	THR	25	-69.090	79.338	-8.738	1.00	20.00
	ATOM	185	OG1	THR	25	-70.120	79.769	-7.852	1.00	20.00
55	ATOM	186	CG2	THR	25	-69.433	77.975	-9.315	1.00	20.00
	ATOM	187	N	PRO	26	-65.938	80.205	-9.249	1.00	84.19
	ATOM	188	CA	PRO	26	-64.567	80.293	-9.904	1.00	82.44
	ATOM	189	C	PRO	26	-63.381	80.037	-8.957	1.00	84.32
	ATOM	190	O	PRO	26	-62.227	79.868	-9.357	1.00	81.24
60	ATOM	191	CB	PRO	26	-64.536	81.691	-10.480	1.00	78.00
	ATOM	192	CG	PRO	26	-65.933	81.872	-10.965	1.00	20.00
	ATOM	193	CD	PRO	26	-66.801	80.898	-10.203	1.00	20.00
	ATOM	194	N	ARG	27	-63.720	80.034	-7.645	1.00	87.06
	ATOM	195	CA	ARG	27	-62.847	79.967	-6.441	1.00	88.17
65	ATOM	196	C	ARG	27	-61.717	81.019	-6.630	1.00	88.45
	ATOM	197	O	ARG	27	-60.619	80.899	-6.089	1.00	84.89

	ATOM	198	CB	ARG	27	-62.558	78.490	-6.004	0.00	86.84
	ATOM	199	CG	ARG	27	-63.782	77.815	-5.264	0.00	85.55
	ATOM	200	CD	ARG	27	-63.561	77.083	-3.883	1.00	20.00
	ATOM	201	NE	ARG	27	-64.782	76.486	-3.267	1.00	20.00
5	ATOM	202	CZ	ARG	27	-64.890	75.834	-2.100	1.00	20.00
	ATOM	203	NH1	ARG	27	-63.819	75.670	-1.351	1.00	20.00
	ATOM	204	NH2	ARG	27	-66.060	75.348	-1.713	1.00	20.00
	ATOM	205	N	PRO	28	-62.100	82.040	-7.423	1.00	89.49
	ATOM	206	CA	PRO	28	-61.252	83.275	-7.550	1.00	91.86
10	ATOM	207	C	PRO	28	-60.984	83.902	-6.136	1.00	95.75
	ATOM	208	O	PRO	28	-60.176	84.811	-5.958	1.00	96.59
	ATOM	209	CB	PRO	28	-61.986	84.093	-8.619	0.00	91.23
	ATOM	210	CG	PRO	28	-62.445	83.005	-9.559	1.00	20.00
	ATOM	211	CD	PRO	28	-62.554	81.723	-8.783	1.00	20.00
15	ATOM	212	N	LYS	29	-61.719	83.349	-5.142	1.00	98.41
	ATOM	213	CA	LYS	29	-61.754	83.674	-3.699	1.00	96.39
	ATOM	214	C	LYS	29	-61.292	85.020	-3.154	1.00	96.88
	ATOM	215	O	LYS	29	-60.097	85.301	-3.105	1.00	94.52
	ATOM	216	CB	LYS	29	-61.012	82.569	-2.949	0.00	95.29
20	ATOM	217	CG	LYS	29	-60.070	81.763	-3.818	0.00	92.34
	ATOM	218	CD	LYS	29	-58.631	82.185	-3.606	0.00	89.50
	ATOM	219	CE	LYS	29	-57.665	81.064	-3.967	1.00	20.00
	ATOM	220	NZ	LYS	29	-56.612	80.882	-2.932	1.00	20.00
	ATOM	221	N	PRO	30	-62.292	85.892	-2.773	1.00	97.72
25	ATOM	222	CA	PRO	30	-61.980	87.142	-1.998	1.00	98.94
	ATOM	223	C	PRO	30	-61.468	86.917	-0.571	1.00	100.71
	ATOM	224	O	PRO	30	-62.157	86.281	0.238	1.00	100.56
	ATOM	225	CB	PRO	30	-63.251	87.948	-2.056	0.00	97.33
	ATOM	226	CG	PRO	30	-63.746	87.672	-3.424	1.00	20.00
30	ATOM	227	CD	PRO	30	-63.120	86.376	-3.881	1.00	20.00
	ATOM	228	N	ALA	31	-60.281	87.431	-0.244	1.00	101.17
	ATOM	229	CA	ALA	31	-59.734	87.261	1.082	1.00	99.22
	ATOM	230	C	ALA	31	-59.821	88.556	1.886	1.00	99.01
	ATOM	231	O	ALA	31	-59.699	88.560	3.113	1.00	102.17
35	ATOM	232	CB	ALA	31	-58.295	86.777	0.984	1.00	20.00
	ATOM	233	N	GLY	32	-60.070	89.628	1.157	1.00	95.52
	ATOM	234	CA	GLY	32	-60.303	90.877	1.793	1.00	91.44
	ATOM	235	C	GLY	32	-61.813	91.032	1.870	1.00	88.49
	ATOM	236	O	GLY	32	-62.400	91.770	1.094	1.00	86.70
40	ATOM	237	N	ALA	33	-62.431	90.344	2.819	1.00	15.00
	ATOM	238	CA	ALA	33	-63.864	90.452	3.014	1.00	15.00
	ATOM	239	C	ALA	33	-64.686	89.314	2.458	1.00	15.00
	ATOM	240	O	ALA	33	-65.746	88.979	2.992	1.00	15.00
	ATOM	241	CB	ALA	33	-64.358	91.767	2.414	1.00	15.00
45	ATOM	242	N	SER	34	-64.221	88.701	1.402	1.00	15.00
	ATOM	243	CA	SER	34	-65.032	87.664	0.761	1.00	15.00
	ATOM	244	C	SER	34	-65.001	86.302	1.448	1.00	15.00
	ATOM	245	O	SER	34	-65.993	85.570	1.457	1.00	15.00
	ATOM	246	CB	SER	34	-64.585	87.512	-0.685	1.00	15.00
50	ATOM	247	OG	SER	34	-65.596	86.888	-1.458	1.00	20.00
	ATOM	248	N	SER	35	-63.869	85.945	2.026	1.00	15.00
	ATOM	249	CA	SER	35	-63.707	84.641	2.681	1.00	15.00
	ATOM	250	C	SER	35	-64.936	84.337	3.538	1.00	15.00
	ATOM	251	O	SER	35	-65.436	83.215	3.614	1.00	15.00
55	ATOM	252	CB	SER	35	-62.428	84.607	3.534	1.00	15.00
	ATOM	253	OG	SER	35	-61.630	85.758	3.298	1.00	20.00
	ATOM	254	N	PRO	36	-65.383	85.414	4.174	1.00	15.00
	ATOM	255	CA	PRO	36	-66.600	85.391	5.032	1.00	15.00
	ATOM	256	C	PRO	36	-67.787	84.746	4.370	1.00	15.00
60	ATOM	257	O	PRO	36	-68.390	83.814	4.903	1.00	15.00
	ATOM	258	CB	PRO	36	-66.677	86.840	5.514	1.00	15.00
	ATOM	259	CG	PRO	36	-65.222	87.191	5.697	1.00	15.00
	ATOM	260	CD	PRO	36	-64.415	86.336	4.783	1.00	15.00
	ATOM	261	N	ALA	37	-68.105	85.245	3.190	1.00	15.00
65	ATOM	262	CA	ALA	37	-69.180	84.704	2.385	1.00	15.00
	ATOM	263	C	ALA	37	-68.864	83.261	2.025	1.00	15.00

	ATOM	264	O	ALA	37	-69.752	82.421	1.955	1.00	15.00
	ATOM	265	CB	ALA	37	-69.372	85.539	1.134	1.00	15.00
	ATOM	266	N	PRO	38	-67.565	82.988	1.748	1.00	15.00
	ATOM	267	CA	PRO	38	-67.137	81.602	1.371	1.00	15.00
5	ATOM	268	C	PRO	38	-67.579	80.516	2.275	1.00	15.00
	ATOM	269	O	PRO	38	-68.176	79.527	1.873	1.00	15.00
	ATOM	270	CB	PRO	38	-65.653	81.696	1.205	1.00	15.00
	ATOM	271	CG	PRO	38	-65.543	83.034	0.533	1.00	20.00
	ATOM	272	CD	PRO	38	-66.742	83.870	0.923	1.00	20.00
10	ATOM	273	N	ARG	39	-67.275	80.741	3.546	1.00	15.00
	ATOM	274	CA	ARG	39	-67.617	79.771	4.585	1.00	15.00
	ATOM	275	C	ARG	39	-69.090	79.907	5.026	1.00	15.00
	ATOM	276	O	ARG	39	-69.530	79.155	5.900	1.00	15.00
	ATOM	277	CB	ARG	39	-66.713	79.908	5.789	1.00	15.00
15	ATOM	278	CG	ARG	39	-65.244	79.885	5.468	1.00	15.00
	ATOM	279	CD	ARG	39	-64.543	80.900	6.340	1.00	15.00
	ATOM	280	NE	ARG	39	-63.113	80.911	6.117	1.00	20.00
	ATOM	281	CZ	ARG	39	-62.224	81.185	7.065	1.00	20.00
	ATOM	282	NH1	ARG	39	-62.632	81.444	8.300	1.00	20.00
20	ATOM	283	NH2	ARG	39	-60.930	81.199	6.778	1.00	20.00
	ATOM	284	N	THR	40	-69.815	80.873	4.456	1.00	15.00
	ATOM	285	CA	THR	40	-71.236	81.187	4.758	1.00	15.00
	ATOM	286	C	THR	40	-72.146	80.307	3.895	1.00	15.00
	ATOM	287	O	THR	40	-73.085	79.656	4.341	1.00	15.00
25	ATOM	288	CB	THR	40	-71.555	82.632	4.454	1.00	15.00
	ATOM	289	OG1	THR	40	-70.505	83.449	4.997	1.00	20.00
	ATOM	290	CG2	THR	40	-72.883	83.028	5.069	1.00	20.00
	ATOM	291	N	ALA	41	-71.793	80.327	2.613	1.00	15.00
	ATOM	292	CA	ALA	41	-72.483	79.507	1.656	1.00	15.00
30	ATOM	293	C	ALA	41	-71.969	78.071	1.722	1.00	15.00
	ATOM	294	O	ALA	41	-72.613	77.173	1.210	1.00	15.00
	ATOM	295	CB	ALA	41	-72.314	80.045	0.240	1.00	15.00
	ATOM	296	N	LEU	42	-70.830	77.856	2.321	1.00	15.00
	ATOM	297	CA	LEU	42	-70.272	76.516	2.455	1.00	15.00
35	ATOM	298	C	LEU	42	-70.967	75.794	3.599	1.00	15.00
	ATOM	299	O	LEU	42	-71.374	74.635	3.470	1.00	15.00
	ATOM	300	CB	LEU	42	-68.777	76.570	2.722	1.00	15.00
	ATOM	301	CG	LEU	42	-67.889	77.070	1.582	1.00	20.00
	ATOM	302	CD1	LEU	42	-66.420	77.039	1.991	1.00	20.00
40	ATOM	303	CD2	LEU	42	-68.114	76.251	0.328	1.00	20.00
	ATOM	304	N	GLN	43	-71.118	76.478	4.722	1.00	15.00
	ATOM	305	CA	GLN	43	-71.825	75.944	5.898	1.00	15.00
	ATOM	306	C	GLN	43	-73.239	75.565	5.529	1.00	15.00
	ATOM	307	O	GLN	43	-73.629	74.402	5.606	1.00	15.00
45	ATOM	308	CB	GLN	43	-71.865	76.976	7.044	1.00	15.00
	ATOM	309	CG	GLN	43	-70.563	77.669	7.370	1.00	20.00
	ATOM	310	CD	GLN	43	-70.795	78.715	8.442	1.00	20.00
	ATOM	311	OE1	GLN	43	-71.922	78.981	8.849	1.00	20.00
	ATOM	312	NE2	GLN	43	-69.845	79.418	9.052	1.00	20.00
50	END					-94.358	153.908	54.620	0.00	0.00

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	ATOM	2	CA	PRO	1	-78.512	94.622	-46.391	1.00	98.53
	ATOM	3	C	PRO	1	-78.084	93.530	-47.292	1.00	101.54
5	ATOM	4	O	PRO	1	-78.291	93.696	-48.503	1.00	102.94
	ATOM	5	CB	PRO	1	-79.505	95.551	-47.062	1.00	96.91
	ATOM	6	CG	PRO	1	-79.433	96.720	-46.147	1.00	20.00
	ATOM	7	CD	PRO	1	-78.123	96.664	-45.378	1.00	20.00
	ATOM	8	N	THR	2	-77.507	92.450	-46.906	1.00	104.82
10	ATOM	9	CA	THR	2	-77.320	91.621	-48.052	1.00	107.58
	ATOM	10	C	THR	2	-78.341	90.500	-47.961	1.00	108.57
	ATOM	11	O	THR	2	-79.389	90.723	-47.354	1.00	109.82
	ATOM	12	CB	THR	2	-75.870	91.194	-48.197	1.00	20.00
	ATOM	13	OG1	THR	2	-75.144	92.197	-48.941	1.00	20.00
15	ATOM	14	CG2	THR	2	-75.789	89.850	-48.902	1.00	20.00
	ATOM	15	N	PRO	3	-78.140	89.288	-48.528	1.00	108.52
	ATOM	16	CA	PRO	3	-79.306	88.363	-48.521	1.00	109.56
	ATOM	17	C	PRO	3	-80.059	88.160	-47.250	1.00	110.79
	ATOM	18	O	PRO	3	-79.449	87.802	-46.242	1.00	114.50
20	ATOM	19	CB	PRO	3	-78.758	87.089	-49.131	1.00	108.23
	ATOM	20	CG	PRO	3	-77.880	87.653	-50.196	1.00	109.80
	ATOM	21	CD	PRO	3	-77.467	89.057	-49.800	1.00	110.35
	ATOM	22	C	CYS	4	-83.124	87.470	-45.611	1.00	101.54
	ATOM	23	O	CYS	4	-84.163	88.117	-45.792	1.00	102.94
25	ATOM	24	CB	CYS	4	-81.410	89.175	-44.960	1.00	96.91
	ATOM	25	SG	CYS	4	-80.133	88.799	-43.717	1.00	92.39
	ATOM	26	N	CYS	4	-81.362	88.324	-47.207	1.00	93.44
	ATOM	27	CA	CYS	4	-81.733	88.016	-45.832	1.00	98.53
	ATOM	28	N	VAL	5	-83.044	86.239	-45.165	1.00	104.82
30	ATOM	29	CA	VAL	5	-84.173	85.388	-44.887	1.00	107.58
	ATOM	30	C	VAL	5	-85.118	85.967	-43.796	1.00	108.57
	ATOM	31	O	VAL	5	-84.648	86.545	-42.811	1.00	109.82
	ATOM	32	CB	VAL	5	-83.578	83.988	-44.668	1.00	109.37
	ATOM	33	CG1	VAL	5	-84.674	82.933	-44.711	1.00	20.00
35	ATOM	34	CG2	VAL	5	-82.492	83.684	-45.684	1.00	20.00
	ATOM	35	N	PRO	6	-86.443	85.812	-44.003	1.00	108.52
	ATOM	36	CA	PRO	6	-87.351	86.408	-42.996	1.00	109.56
	ATOM	37	C	PRO	6	-87.001	86.104	-41.532	1.00	110.79
	ATOM	38	O	PRO	6	-86.658	84.975	-41.207	1.00	114.50
40	ATOM	39	CB	PRO	6	-88.732	86.027	-43.511	1.00	108.23
	ATOM	40	CG	PRO	6	-88.587	86.160	-44.982	1.00	109.80
	ATOM	41	CD	PRO	6	-87.120	85.985	-45.310	1.00	110.35
	ATOM	42	N	ALA	7	-87.102	87.089	-40.644	1.00	109.87
	ATOM	43	CA	ALA	7	-86.721	86.944	-39.239	1.00	107.78
45	ATOM	44	C	ALA	7	-85.288	87.435	-38.955	1.00	105.15
	ATOM	45	O	ALA	7	-85.093	88.119	-37.965	1.00	103.42
	ATOM	46	CB	ALA	7	-86.872	85.503	-38.797	1.00	109.08
	ATOM	47	N	GLU	8	-84.257	87.159	-39.817	1.00	101.11
	ATOM	48	CA	GLU	8	-82.840	87.567	-39.456	1.00	95.31
50	ATOM	49	CB	GLU	8	-81.827	86.509	-39.808	1.00	92.47
	ATOM	50	CG	GLU	8	-82.424	85.188	-40.255	1.00	86.82
	ATOM	51	CD	GLU	8	-81.753	84.660	-41.512	1.00	85.55
	ATOM	52	OE1	GLU	8	-81.700	85.399	-42.521	1.00	85.72
	ATOM	53	OE2	GLU	8	-81.266	83.520	-41.482	1.00	81.01
55	ATOM	54	C	GLU	8	-82.299	88.874	-40.005	1.00	92.66
	ATOM	55	O	GLU	8	-82.842	89.451	-40.955	1.00	94.43
	ATOM	56	N	CYS	9	-81.210	89.331	-39.396	1.00	85.19
	ATOM	57	CA	CYS	9	-80.592	90.591	-39.773	1.00	78.14
	ATOM	58	C	CYS	9	-79.123	90.348	-40.055	1.00	75.43
60	ATOM	59	O	CYS	9	-78.452	89.624	-39.325	1.00	76.93
	ATOM	60	CB	CYS	9	-80.747	91.628	-38.662	1.00	76.59
	ATOM	61	SG	CYS	9	-79.796	91.269	-37.151	1.00	20.00
	ATOM	62	N	PHE	10	-78.654	90.973	-41.133	1.00	69.11
	ATOM	63	CA	PHE	10	-77.238	90.916	-41.569	1.00	62.54
65	ATOM	64	CB	PHE	10	-77.063	91.634	-42.963	1.00	61.14
	ATOM	65	CG	PHE	10	-75.667	91.566	-43.542	1.00	61.11

	ATOM	66	CD1	PHE	10	-75.192	90.348	-44.017	1.00	59.90
	ATOM	67	CD2	PHE	10	-74.825	92.675	-43.573	1.00	63.23
	ATOM	68	CE1	PHE	10	-73.893	90.237	-44.514	1.00	61.45
	ATOM	69	CE2	PHE	10	-73.518	92.583	-44.068	1.00	61.15
5	ATOM	70	CZ	PHE	10	-73.050	91.364	-44.538	1.00	61.89
	ATOM	71	C	PHE	10	-76.339	91.578	-40.510	1.00	60.79
	ATOM	72	O	PHE	10	-76.470	92.771	-40.220	1.00	63.10
	ATOM	73	N	ASP	11	-75.431	90.801	-39.904	1.00	57.06
	ATOM	74	CA	ASP	11	-74.497	91.361	-38.904	1.00	52.20
10	ATOM	75	CB	ASP	11	-74.136	90.295	-37.802	1.00	46.52
	ATOM	76	CG	ASP	11	-73.255	90.787	-36.639	1.00	47.00
	ATOM	77	OD1	ASP	11	-72.416	91.682	-36.884	1.00	49.00
	ATOM	78	OD2	ASP	11	-73.424	90.282	-35.511	1.00	48.43
	ATOM	79	C	ASP	11	-73.221	91.792	-39.656	1.00	53.02
15	ATOM	80	O	ASP	11	-72.567	90.941	-40.251	1.00	53.25
	ATOM	81	N	LEU	12	-72.854	93.086	-39.642	1.00	55.34
	ATOM	82	CA	LEU	12	-71.644	93.586	-40.335	1.00	54.13
	ATOM	83	C	LEU	12	-70.330	93.194	-39.636	1.00	54.96
	ATOM	84	O	LEU	12	-69.250	93.227	-40.220	1.00	54.11
20	ATOM	85	CB	LEU	12	-71.745	95.113	-40.533	1.00	54.65
	ATOM	86	CG	LEU	12	-73.039	95.673	-41.149	1.00	20.00
	ATOM	87	CD1	LEU	12	-72.997	97.192	-41.164	1.00	20.00
	ATOM	88	CD2	LEU	12	-73.233	95.116	-42.551	1.00	20.00
	ATOM	89	N	LEU	13	-70.441	92.807	-38.351	1.00	49.64
25	ATOM	90	CA	LEU	13	-69.289	92.385	-37.564	1.00	46.11
	ATOM	91	CB	LEU	13	-69.438	92.686	-36.099	1.00	40.39
	ATOM	92	CG	LEU	13	-68.229	92.128	-35.382	1.00	34.58
	ATOM	93	CD1	LEU	13	-67.116	93.170	-35.330	1.00	33.04
	ATOM	94	CD2	LEU	13	-68.591	91.655	-33.989	1.00	33.82
30	ATOM	95	C	LEU	13	-69.028	90.894	-37.709	1.00	49.82
	ATOM	96	O	LEU	13	-68.041	90.400	-37.168	1.00	50.94
	ATOM	97	N	VAL	14	-69.883	90.164	-38.430	1.00	51.20
	ATOM	98	CA	VAL	14	-69.481	88.774	-38.725	1.00	51.39
	ATOM	99	C	VAL	14	-69.405	88.634	-40.211	1.00	53.87
35	ATOM	100	O	VAL	14	-68.793	87.706	-40.744	1.00	50.27
	ATOM	101	CB	VAL	14	-70.441	87.659	-38.352	1.00	47.82
	ATOM	102	CG1	VAL	14	-69.811	86.302	-38.653	1.00	20.00
	ATOM	103	CG2	VAL	14	-70.857	87.749	-36.889	1.00	20.00
	ATOM	104	N	ARG	15	-70.088	89.577	-40.875	1.00	58.57
40	ATOM	105	CA	ARG	15	-70.168	89.711	-42.310	1.00	64.33
	ATOM	106	C	ARG	15	-70.979	88.573	-42.941	1.00	66.69
	ATOM	107	O	ARG	15	-70.718	88.113	-44.063	1.00	67.11
	ATOM	108	CB	ARG	15	-68.725	89.900	-42.839	1.00	65.68
	ATOM	109	CG	ARG	15	-67.869	90.927	-42.076	1.00	69.20
45	ATOM	110	CD	ARG	15	-67.169	91.925	-42.994	1.00	20.00
	ATOM	111	NE	ARG	15	-66.186	92.801	-42.313	1.00	20.00
	ATOM	112	CZ	ARG	15	-65.453	93.717	-42.937	1.00	20.00
	ATOM	113	NH1	ARG	15	-65.568	93.873	-44.249	1.00	20.00
	ATOM	114	NH2	ARG	15	-64.610	94.473	-42.248	1.00	20.00
50	ATOM	115	N	HIS	16	-71.979	88.139	-42.149	1.00	69.57
	ATOM	116	CA	HIS	16	-72.907	87.084	-42.517	1.00	71.62
	ATOM	117	C	HIS	16	-74.235	87.170	-41.830	1.00	73.99
	ATOM	118	O	HIS	16	-74.383	87.831	-40.810	1.00	75.45
	ATOM	119	CB	HIS	16	-72.444	85.717	-42.085	1.00	69.66
55	ATOM	120	CG	HIS	16	-71.058	85.378	-42.520	1.00	20.00
	ATOM	121	ND1	HIS	16	-70.816	84.878	-43.782	1.00	20.00
	ATOM	122	CD2	HIS	16	-69.861	85.452	-41.886	1.00	20.00
	ATOM	123	CE1	HIS	16	-69.528	84.662	-43.906	1.00	20.00
	ATOM	124	NE2	HIS	16	-68.925	84.985	-42.776	1.00	20.00
60	ATOM	125	N	CYS	17	-75.202	86.477	-42.389	1.00	75.75
	ATOM	126	CA	CYS	17	-76.521	86.534	-41.802	1.00	77.01
	ATOM	127	C	CYS	17	-76.582	85.806	-40.446	1.00	75.86
	ATOM	128	O	CYS	17	-75.886	84.826	-40.198	1.00	73.08
	ATOM	129	CB	CYS	17	-77.547	85.961	-42.777	1.00	80.96
65	ATOM	130	SG	CYS	17	-78.866	87.132	-43.247	1.00	86.58
	ATOM	131	N	VAL	18	-77.447	86.334	-39.580	1.00	75.89

	ATOM	132	CA	VAL	18	-77.674	85.821	-38.258	1.00	77.02
	ATOM	133	C	VAL	18	-79.151	85.779	-38.023	1.00	78.19
	ATOM	134	O	VAL	18	-79.921	86.523	-38.609	1.00	77.54
	ATOM	135	CB	VAL	18	-77.017	86.661	-37.131	1.00	76.68
5	ATOM	136	CG1	VAL	18	-77.861	86.628	-35.874	1.00	76.28
	ATOM	137	CG2	VAL	18	-75.626	86.141	-36.833	1.00	76.37
	ATOM	138	N	ALA	19	-79.531	84.888	-37.104	1.00	80.09
	ATOM	139	CA	ALA	19	-80.951	84.682	-36.668	1.00	82.16
	ATOM	140	C	ALA	19	-81.309	85.599	-35.531	1.00	84.28
10	ATOM	141	O	ALA	19	-80.552	85.633	-34.560	1.00	84.12
	ATOM	142	CB	ALA	19	-81.174	83.224	-36.261	1.00	80.00
	ATOM	143	N	CYS	20	-82.354	86.380	-35.561	1.00	87.37
	ATOM	144	CA	CYS	20	-82.379	87.319	-34.422	1.00	88.33
	ATOM	145	CB	CYS	20	-83.409	88.426	-34.678	1.00	86.90
15	ATOM	146	SG	CYS	20	-82.700	89.999	-35.260	1.00	84.56
	ATOM	147	C	CYS	20	-82.538	86.653	-33.054	1.00	89.64
	ATOM	148	O	CYS	20	-82.555	87.331	-32.031	1.00	89.77
	ATOM	149	N	GLY	21	-82.671	85.328	-33.037	1.00	87.37
	ATOM	150	CA	GLY	21	-82.836	84.656	-31.770	1.00	88.33
20	ATOM	151	C	GLY	21	-81.584	84.802	-30.904	1.00	89.64
	ATOM	152	O	GLY	21	-81.571	84.518	-29.709	1.00	89.77
	ATOM	153	N	LEU	22	-80.488	85.245	-31.531	1.00	87.56
	ATOM	154	CA	LEU	22	-79.199	85.410	-30.857	1.00	85.33
	ATOM	155	CB	LEU	22	-78.052	85.288	-31.843	1.00	79.98
25	ATOM	156	CG	LEU	22	-77.157	84.069	-31.669	1.00	77.91
	ATOM	157	CD1	LEU	22	-76.306	83.838	-32.903	1.00	78.13
	ATOM	158	CD2	LEU	22	-76.279	84.225	-30.427	1.00	72.50
	ATOM	159	C	LEU	22	-79.083	86.735	-30.126	1.00	86.05
	ATOM	160	O	LEU	22	-78.685	86.747	-28.968	1.00	88.06
30	ATOM	161	N	LEU	23	-79.418	87.861	-30.804	1.00	85.11
	ATOM	162	CA	LEU	23	-79.231	89.172	-30.173	1.00	83.33
	ATOM	163	C	LEU	23	-80.363	89.629	-29.256	1.00	84.21
	ATOM	164	O	LEU	23	-80.263	90.717	-28.666	1.00	86.28
	ATOM	165	CB	LEU	23	-79.001	90.248	-31.240	1.00	78.93
35	ATOM	166	CG	LEU	23	-78.079	89.997	-32.423	1.00	73.68
	ATOM	167	CD1	LEU	23	-78.375	90.999	-33.518	1.00	20.00
	ATOM	168	CD2	LEU	23	-76.627	90.043	-32.014	1.00	20.00
	ATOM	169	N	ARG	24	-81.414	88.842	-29.091	1.00	86.09
	ATOM	170	CA	ARG	24	-82.487	89.356	-28.236	1.00	85.39
40	ATOM	171	C	ARG	24	-82.374	88.908	-26.797	1.00	84.10
	ATOM	172	O	ARG	24	-82.889	89.581	-25.906	1.00	83.24
	ATOM	173	CB	ARG	24	-83.845	89.029	-28.790	1.00	80.05
	ATOM	174	CG	ARG	24	-83.943	87.758	-29.581	1.00	20.00
	ATOM	175	CD	ARG	24	-85.388	87.347	-29.632	1.00	20.00
45	ATOM	176	NE	ARG	24	-85.960	87.601	-30.928	1.00	20.00
	ATOM	177	CZ	ARG	24	-87.143	87.096	-31.236	1.00	20.00
	ATOM	178	NH1	ARG	24	-87.809	86.372	-30.346	1.00	20.00
	ATOM	179	NH2	ARG	24	-87.658	87.315	-32.428	1.00	20.00
	ATOM	180	N	THR	25	-81.719	87.761	-26.528	1.00	84.61
50	ATOM	181	CA	THR	25	-81.572	87.288	-25.132	1.00	85.07
	ATOM	182	C	THR	25	-80.478	88.116	-24.458	1.00	85.65
	ATOM	183	O	THR	25	-80.488	88.367	-23.259	1.00	86.77
	ATOM	184	CB	THR	25	-81.319	85.768	-25.111	1.00	20.00
	ATOM	185	OG1	THR	25	-82.284	85.127	-25.944	1.00	20.00
55	ATOM	186	CG2	THR	25	-81.419	85.236	-23.692	1.00	20.00
	ATOM	187	N	PRO	26	-79.507	88.526	-25.337	1.00	84.19
	ATOM	188	CA	PRO	26	-78.445	89.569	-25.021	1.00	82.44
	ATOM	189	C	PRO	26	-78.963	91.017	-24.941	1.00	84.32
	ATOM	190	O	PRO	26	-78.289	91.949	-24.498	1.00	81.24
60	ATOM	191	CB	PRO	26	-77.432	89.395	-26.130	1.00	78.00
	ATOM	192	CG	PRO	26	-77.413	87.917	-26.321	1.00	20.00
	ATOM	193	CD	PRO	26	-78.717	87.380	-25.779	1.00	20.00
	ATOM	194	N	ARG	27	-80.222	91.173	-25.419	1.00	87.06
	ATOM	195	CA	ARG	27	-81.002	92.421	-25.643	1.00	88.17
65	ATOM	196	C	ARG	27	-80.082	93.410	-26.414	1.00	88.45
	ATOM	197	O	ARG	27	-80.228	94.628	-26.338	1.00	84.89

	ATOM	198	CB	ARG	27	-81.815	92.844	-24.372	0.00	86.84
	ATOM	199	CG	ARG	27	-83.112	91.966	-24.148	0.00	85.55
	ATOM	200	CD	ARG	27	-84.503	92.669	-23.895	1.00	20.00
	ATOM	201	NE	ARG	27	-85.662	91.750	-23.703	1.00	20.00
5	ATOM	202	CZ	ARG	27	-86.949	92.067	-23.497	1.00	20.00
	ATOM	203	NH1	ARG	27	-87.301	93.336	-23.452	1.00	20.00
	ATOM	204	NH2	ARG	27	-87.852	91.114	-23.326	1.00	20.00
	ATOM	205	N	PRO	28	-79.154	92.770	-27.154	1.00	89.49
	ATOM	206	CA	PRO	28	-78.317	93.518	-28.154	1.00	91.86
10	ATOM	207	C	PRO	28	-79.234	94.280	-29.174	1.00	95.75
	ATOM	208	O	PRO	28	-78.794	95.100	-29.976	1.00	96.59
	ATOM	209	CB	PRO	28	-77.339	92.451	-28.658	0.00	91.23
	ATOM	210	CG	PRO	28	-77.063	91.680	-27.389	1.00	20.00
	ATOM	211	CD	PRO	28	-78.234	91.855	-26.464	1.00	20.00
15	ATOM	212	N	LYS	29	-80.546	93.950	-29.086	1.00	98.41
	ATOM	213	CA	LYS	29	-81.698	94.438	-29.873	1.00	96.39
	ATOM	214	C	LYS	29	-81.537	95.070	-31.251	1.00	96.88
	ATOM	215	O	LYS	29	-81.079	96.204	-31.373	1.00	94.52
	ATOM	216	CB	LYS	29	-82.498	95.398	-28.994	0.00	95.29
20	ATOM	217	CG	LYS	29	-81.713	95.961	-27.828	0.00	92.34
	ATOM	218	CD	LYS	29	-81.264	97.381	-28.104	0.00	89.50
	ATOM	219	CE	LYS	29	-81.027	98.149	-26.810	1.00	20.00
	ATOM	220	NZ	LYS	29	-81.640	99.503	-26.847	1.00	20.00
	ATOM	221	N	PRO	30	-81.893	94.276	-32.324	1.00	97.72
25	ATOM	222	CA	PRO	30	-82.016	94.850	-33.708	1.00	98.94
	ATOM	223	C	PRO	30	-83.169	95.842	-33.905	1.00	100.71
	ATOM	224	O	PRO	30	-84.332	95.488	-33.671	1.00	100.56
	ATOM	225	CB	PRO	30	-82.104	93.646	-34.608	0.00	97.33
	ATOM	226	CG	PRO	30	-81.176	92.691	-33.960	1.00	20.00
30	ATOM	227	CD	PRO	30	-81.032	93.106	-32.515	1.00	20.00
	ATOM	228	N	ALA	31	-82.871	97.068	-34.339	1.00	101.17
	ATOM	229	CA	ALA	31	-83.904	98.055	-34.549	1.00	99.22
	ATOM	230	C	ALA	31	-84.170	98.267	-36.038	1.00	99.01
	ATOM	231	O	ALA	31	-85.197	98.824	-36.433	1.00	102.17
35	ATOM	232	CB	ALA	31	-83.509	99.361	-33.876	1.00	20.00
	ATOM	233	N	GLY	32	-83.235	97.776	-36.829	1.00	95.52
	ATOM	234	CA	GLY	32	-83.421	97.790	-38.237	1.00	91.44
	ATOM	235	C	GLY	32	-83.939	96.411	-38.606	1.00	88.49
	ATOM	236	O	GLY	32	-83.196	95.586	-39.114	1.00	86.70
40	ATOM	237	N	ALA	33	-85.219	96.174	-38.360	1.00	15.00
	ATOM	238	CA	ALA	33	-85.832	94.908	-38.714	1.00	15.00
	ATOM	239	C	ALA	33	-86.030	93.938	-37.575	1.00	15.00
	ATOM	240	O	ALA	33	-86.970	93.140	-37.579	1.00	15.00
	ATOM	241	CB	ALA	33	-85.004	94.234	-39.807	1.00	15.00
45	ATOM	242	N	SER	34	-85.173	93.989	-36.590	1.00	15.00
	ATOM	243	CA	SER	34	-85.257	92.998	-35.514	1.00	15.00
	ATOM	244	C	SER	34	-86.333	93.271	-34.468	1.00	15.00
	ATOM	245	O	SER	34	-86.936	92.347	-33.917	1.00	15.00
	ATOM	246	CB	SER	34	-83.902	92.895	-34.832	1.00	15.00
50	ATOM	247	OG	SER	34	-83.790	91.671	-34.125	1.00	20.00
	ATOM	248	N	SER	35	-86.584	94.535	-34.178	1.00	15.00
	ATOM	249	CA	SER	35	-87.568	94.918	-33.157	1.00	15.00
	ATOM	250	C	SER	35	-88.836	94.080	-33.319	1.00	15.00
	ATOM	251	O	SER	35	-89.471	93.637	-32.362	1.00	15.00
55	ATOM	252	CB	SER	35	-87.895	96.418	-33.241	1.00	15.00
	ATOM	253	OG	SER	35	-87.009	97.081	-34.132	1.00	20.00
	ATOM	254	N	PRO	36	-89.155	93.894	-34.596	1.00	15.00
	ATOM	255	CA	PRO	36	-90.318	93.068	-35.019	1.00	15.00
	ATOM	256	C	PRO	36	-90.371	91.720	-34.352	1.00	15.00
60	ATOM	257	O	PRO	36	-91.371	91.348	-33.739	1.00	15.00
	ATOM	258	CB	PRO	36	-90.246	93.175	-36.543	1.00	15.00
	ATOM	259	CG	PRO	36	-89.792	94.599	-36.740	1.00	15.00
	ATOM	260	CD	PRO	36	-89.030	95.020	-35.532	1.00	15.00
	ATOM	261	N	ALA	37	-89.271	91.000	-34.467	1.00	15.00
65	ATOM	262	CA	ALA	37	-89.123	89.706	-33.836	1.00	15.00
	ATOM	263	C	ALA	37	-89.220	89.867	-32.327	1.00	15.00

	ATOM	264	O	ALA	37	-89.758	89.012	-31.636	1.00	15.00
	ATOM	265	CB	ALA	37	-87.799	89.079	-34.224	1.00	15.00
	ATOM	266	N	PRO	38	-88.640	90.978	-31.809	1.00	15.00
	ATOM	267	CA	PRO	38	-88.665	91.238	-30.333	1.00	15.00
5	ATOM	268	C	PRO	38	-89.989	91.149	-29.678	1.00	15.00
	ATOM	269	O	PRO	38	-90.193	90.444	-28.699	1.00	15.00
	ATOM	270	CB	PRO	38	-87.988	92.563	-30.171	1.00	15.00
	ATOM	271	CG	PRO	38	-86.887	92.427	-31.182	1.00	20.00
	ATOM	272	CD	PRO	38	-87.330	91.451	-32.250	1.00	20.00
10	ATOM	273	N	ARG	39	-90.914	91.890	-30.271	1.00	15.00
	ATOM	274	CA	ARG	39	-92.281	91.943	-29.755	1.00	15.00
	ATOM	275	C	ARG	39	-93.111	90.728	-30.223	1.00	15.00
	ATOM	276	O	ARG	39	-94.290	90.630	-29.870	1.00	15.00
	ATOM	277	CB	ARG	39	-92.978	93.221	-30.166	1.00	15.00
15	ATOM	278	CG	ARG	39	-92.214	94.475	-29.844	1.00	15.00
	ATOM	279	CD	ARG	39	-92.376	95.446	-30.991	1.00	15.00
	ATOM	280	NE	ARG	39	-91.697	96.699	-30.739	1.00	20.00
	ATOM	281	CZ	ARG	39	-92.127	97.872	-31.194	1.00	20.00
	ATOM	282	NH1	ARG	39	-93.248	97.936	-31.900	1.00	20.00
20	ATOM	283	NH2	ARG	39	-91.438	98.974	-30.941	1.00	20.00
	ATOM	284	N	THR	40	-92.512	89.849	-31.031	1.00	15.00
	ATOM	285	CA	THR	40	-93.139	88.633	-31.612	1.00	15.00
	ATOM	286	C	THR	40	-93.008	87.471	-30.622	1.00	15.00
	ATOM	287	O	THR	40	-93.944	86.754	-30.286	1.00	15.00
25	ATOM	288	CB	THR	40	-92.465	88.230	-32.902	1.00	15.00
	ATOM	289	OG1	THR	40	-92.293	89.407	-33.707	1.00	20.00
	ATOM	290	CG2	THR	40	-93.304	87.214	-33.651	1.00	20.00
	ATOM	291	N	ALA	41	-91.765	87.339	-30.167	1.00	15.00
	ATOM	292	CA	ALA	41	-91.456	86.348	-29.174	1.00	15.00
30	ATOM	293	C	ALA	41	-91.854	86.847	-27.788	1.00	15.00
	ATOM	294	O	ALA	41	-91.946	86.059	-26.864	1.00	15.00
	ATOM	295	CB	ALA	41	-89.972	85.996	-29.183	1.00	15.00
	ATOM	296	N	LEU	42	-92.071	88.124	-27.636	1.00	15.00
	ATOM	297	CA	LEU	42	-92.480	88.690	-26.357	1.00	15.00
35	ATOM	298	C	LEU	42	-93.968	88.452	-26.154	1.00	15.00
	ATOM	299	O	LEU	42	-94.407	88.022	-25.082	1.00	15.00
	ATOM	300	CB	LEU	42	-92.192	90.181	-26.300	1.00	15.00
	ATOM	301	CG	LEU	42	-90.722	90.599	-26.270	1.00	20.00
	ATOM	302	CD1	LEU	42	-90.596	92.117	-26.185	1.00	20.00
40	ATOM	303	CD2	LEU	42	-89.998	89.936	-25.118	1.00	20.00
	ATOM	304	N	GLN	43	-94.753	88.718	-27.187	1.00	15.00
	ATOM	305	CA	GLN	43	-96.206	88.480	-27.172	1.00	15.00
	ATOM	306	C	GLN	43	-96.495	87.027	-26.881	1.00	15.00
	ATOM	307	O	GLN	43	-97.109	86.687	-25.872	1.00	15.00
45	ATOM	308	CB	GLN	43	-96.849	88.858	-28.522	1.00	15.00
	ATOM	309	CG	GLN	43	-96.448	90.192	-29.107	1.00	20.00
	ATOM	310	CD	GLN	43	-97.086	90.365	-30.471	1.00	20.00
	ATOM	311	OE1	GLN	43	-97.724	89.461	-31.003	1.00	20.00
	ATOM	312	NE2	GLN	43	-97.047	91.473	-31.205	1.00	20.00
50	END					-118.331	85.231	-119.150	0.00	0.00

TABLE 32

	ATOM	2	CA	PRO	1	7.426	107.508	31.139	1.00	98.53
	ATOM	3	C	PRO	1	8.876	107.800	31.160	1.00	101.54
5	ATOM	4	O	PRO	1	9.632	106.879	30.820	1.00	102.94
	ATOM	5	CB	PRO	1	7.151	106.017	31.119	1.00	96.91
	ATOM	6	CG	PRO	1	5.729	106.017	30.687	1.00	20.00
	ATOM	7	CD	PRO	1	5.424	107.336	29.995	1.00	20.00
	ATOM	8	N	THR	2	9.387	108.933	31.484	1.00	104.82
10	ATOM	9	CA	THR	2	10.807	108.797	31.449	1.00	107.58
	ATOM	10	C	THR	2	11.283	108.748	32.892	1.00	108.57
	ATOM	11	O	THR	2	10.504	108.317	33.742	1.00	109.82
	ATOM	12	CB	THR	2	11.445	109.858	30.571	1.00	20.00
	ATOM	13	OG1	THR	2	11.478	109.393	29.203	1.00	20.00
15	ATOM	14	CG2	THR	2	12.851	110.165	31.060	1.00	20.00
	ATOM	15	N	PRO	3	12.522	109.149	33.260	1.00	108.52
	ATOM	16	CA	PRO	3	12.909	108.861	34.668	1.00	109.56
	ATOM	17	C	PRO	3	11.962	109.219	35.764	1.00	110.79
	ATOM	18	O	PRO	3	11.551	110.375	35.849	1.00	114.50
20	ATOM	19	CB	PRO	3	14.280	109.491	34.800	1.00	108.23
	ATOM	20	CG	PRO	3	14.862	109.156	33.469	1.00	109.80
	ATOM	21	CD	PRO	3	13.735	108.956	32.476	1.00	110.35
	ATOM	22	C	CYS	4	10.650	108.543	38.986	1.00	101.54
	ATOM	23	O	CYS	4	10.185	107.447	39.324	1.00	102.94
25	ATOM	24	CB	CYS	4	9.367	109.179	36.932	1.00	96.91
	ATOM	25	SG	CYS	4	8.910	110.912	36.607	1.00	92.39
	ATOM	26	N	CYS	4	11.596	108.323	36.653	1.00	93.44
	ATOM	27	CA	CYS	4	10.706	109.039	37.560	1.00	98.53
	ATOM	28	N	VAL	5	11.127	109.465	39.788	1.00	104.82
30	ATOM	29	CA	VAL	5	11.270	109.322	41.215	1.00	107.58
	ATOM	30	C	VAL	5	9.919	109.063	41.940	1.00	108.57
	ATOM	31	O	VAL	5	8.897	109.652	41.577	1.00	109.82
	ATOM	32	CB	VAL	5	12.115	110.528	41.656	1.00	109.37
	ATOM	33	CG1	VAL	5	12.636	110.321	43.071	1.00	20.00
35	ATOM	34	CG2	VAL	5	13.258	110.785	40.690	1.00	20.00
	ATOM	35	N	PRO	6	9.940	108.166	42.949	1.00	108.52
	ATOM	36	CA	PRO	6	8.648	107.872	43.610	1.00	109.56
	ATOM	37	C	PRO	6	7.817	109.102	44.006	1.00	110.79
	ATOM	38	O	PRO	6	8.364	110.071	44.512	1.00	114.50
40	ATOM	39	CB	PRO	6	9.034	106.873	44.690	1.00	108.23
	ATOM	40	CG	PRO	6	10.067	106.042	44.022	1.00	109.80
	ATOM	41	CD	PRO	6	10.682	106.883	42.924	1.00	110.35
	ATOM	42	N	ALA	7	6.503	109.065	43.800	1.00	109.87
	ATOM	43	CA	ALA	7	5.619	110.201	44.061	1.00	107.78
45	ATOM	44	C	ALA	7	5.343	111.045	42.801	1.00	105.15
	ATOM	45	O	ALA	7	4.200	111.408	42.584	1.00	103.42
	ATOM	46	CB	ALA	7	6.194	111.076	45.156	1.00	109.08
	ATOM	47	N	GLU	8	6.343	111.343	41.911	1.00	101.11
	ATOM	48	CA	GLU	8	6.061	112.263	40.736	1.00	95.31
50	ATOM	49	CB	GLU	8	7.183	113.236	40.482	1.00	92.47
	ATOM	50	CG	GLU	8	8.263	113.247	41.547	1.00	86.82
	ATOM	51	CD	GLU	8	9.656	113.208	40.942	1.00	85.55
	ATOM	52	OE1	GLU	8	9.939	112.287	40.142	1.00	85.72
	ATOM	53	OE2	GLU	8	10.454	114.105	41.251	1.00	81.01
55	ATOM	54	C	GLU	8	5.721	111.643	39.395	1.00	92.66
	ATOM	55	O	GLU	8	5.959	110.454	39.152	1.00	94.43
	ATOM	56	N	CYS	9	5.165	112.472	38.518	1.00	85.19
	ATOM	57	CA	CYS	9	4.741	112.025	37.201	1.00	78.14
	ATOM	58	C	CYS	9	5.365	112.927	36.157	1.00	75.43
60	ATOM	59	O	CYS	9	5.407	114.144	36.317	1.00	76.93
	ATOM	60	CB	CYS	9	3.218	112.058	37.079	1.00	76.59
	ATOM	61	SG	CYS	9	2.493	113.729	37.075	1.00	20.00
	ATOM	62	N	PHE	10	5.845	112.291	35.090	1.00	69.11
	ATOM	63	CA	PHE	10	6.455	112.977	33.925	1.00	62.54
65	ATOM	64	CB	PHE	10	7.060	111.921	32.921	1.00	61.14
	ATOM	65	CG	PHE	10	7.780	112.516	31.731	1.00	61.11

	ATOM	66	CD1	PHE	10	9.001	113.149	31.928	1.00	59.90
	ATOM	67	CD2	PHE	10	7.235	112.489	30.448	1.00	63.23
	ATOM	68	CE1	PHE	10	9.671	113.751	30.862	1.00	61.45
	ATOM	69	CE2	PHE	10	7.892	113.088	29.366	1.00	61.15
5	ATOM	70	CZ	PHE	10	9.110	113.720	29.572	1.00	61.89
	ATOM	71	C	PHE	10	5.399	113.844	33.219	1.00	60.79
	ATOM	72	O	PHE	10	4.390	113.336	32.718	1.00	63.10
	ATOM	73	N	ASP	11	5.610	115.167	33.192	1.00	57.06
	ATOM	74	CA	ASP	11	4.669	116.073	32.499	1.00	52.20
10	ATOM	75	CB	ASP	11	4.600	117.476	33.211	1.00	46.52
	ATOM	76	CG	ASP	11	3.573	118.476	32.651	1.00	47.00
	ATOM	77	OD1	ASP	11	3.325	118.430	31.426	1.00	49.00
	ATOM	78	OD2	ASP	11	3.031	119.276	33.441	1.00	48.43
	ATOM	79	C	ASP	11	5.174	116.242	31.052	1.00	53.02
15	ATOM	80	O	ASP	11	6.281	116.738	30.864	1.00	53.25
	ATOM	81	N	LEU	12	4.393	115.844	30.031	1.00	55.34
	ATOM	82	CA	LEU	12	4.798	115.971	28.613	1.00	54.13
	ATOM	83	C	LEU	12	4.763	117.419	28.091	1.00	54.96
	ATOM	84	O	LEU	12	5.365	117.758	27.076	1.00	54.11
20	ATOM	85	CB	LEU	12	3.943	115.033	27.735	1.00	54.65
	ATOM	86	CG	LEU	12	3.812	113.563	28.172	1.00	20.00
	ATOM	87	CD1	LEU	12	2.851	112.828	27.253	1.00	20.00
	ATOM	88	CD2	LEU	12	5.180	112.898	28.180	1.00	20.00
	ATOM	89	N	LEU	13	4.037	118.288	28.820	1.00	49.64
25	ATOM	90	CA	LEU	13	3.927	119.698	28.465	1.00	46.11
	ATOM	91	CB	LEU	13	2.617	120.307	28.882	1.00	40.39
	ATOM	92	CG	LEU	13	2.656	121.779	28.541	1.00	34.58
	ATOM	93	CD1	LEU	13	2.141	122.009	27.124	1.00	33.04
	ATOM	94	CD2	LEU	13	1.862	122.592	29.542	1.00	33.82
30	ATOM	95	C	LEU	13	5.042	120.521	29.089	1.00	49.82
	ATOM	96	O	LEU	13	5.133	121.717	28.819	1.00	50.94
	ATOM	97	N	VAL	14	5.899	119.913	29.914	1.00	51.20
	ATOM	98	CA	VAL	14	7.086	120.690	30.325	1.00	51.39
	ATOM	99	C	VAL	14	8.294	119.942	29.862	1.00	53.87
35	ATOM	100	O	VAL	14	9.397	120.486	29.771	1.00	50.27
	ATOM	101	CB	VAL	14	7.360	120.842	31.810	1.00	47.82
	ATOM	102	CG1	VAL	14	8.569	121.745	32.031	1.00	20.00
	ATOM	103	CG2	VAL	14	6.140	121.382	32.548	1.00	20.00
	ATOM	104	N	ARG	15	8.060	118.648	29.606	1.00	58.57
40	ATOM	105	CA	ARG	15	9.026	117.694	29.116	1.00	64.33
	ATOM	106	C	ARG	15	10.087	117.369	30.175	1.00	66.69
	ATOM	107	O	ARG	15	11.264	117.111	29.880	1.00	67.11
	ATOM	108	CB	ARG	15	9.551	118.221	27.758	1.00	65.68
	ATOM	109	CG	ARG	15	8.473	118.707	26.774	1.00	69.20
45	ATOM	110	CD	ARG	15	8.634	118.127	25.371	1.00	20.00
	ATOM	111	NE	ARG	15	7.735	118.722	24.354	1.00	20.00
	ATOM	112	CZ	ARG	15	7.737	118.375	23.072	1.00	20.00
	ATOM	113	NH1	ARG	15	8.592	117.459	22.635	1.00	20.00
	ATOM	114	NH2	ARG	15	6.885	118.944	22.230	1.00	20.00
50	ATOM	115	N	HIS	16	9.603	117.402	31.431	1.00	69.57
	ATOM	116	CA	HIS	16	10.394	117.113	32.615	1.00	71.62
	ATOM	117	C	HIS	16	9.594	116.616	33.780	1.00	73.99
	ATOM	118	O	HIS	16	8.384	116.788	33.840	1.00	75.45
	ATOM	119	CB	HIS	16	11.037	118.342	33.206	1.00	69.66
55	ATOM	120	CG	HIS	16	11.822	119.148	32.228	1.00	20.00
	ATOM	121	ND1	HIS	16	13.126	118.816	31.924	1.00	20.00
	ATOM	122	CD2	HIS	16	11.513	120.252	31.501	1.00	20.00
	ATOM	123	CE1	HIS	16	13.583	119.681	31.049	1.00	20.00
	ATOM	124	NE2	HIS	16	12.640	120.567	30.783	1.00	20.00
60	ATOM	125	N	CYS	17	10.287	116.012	34.719	1.00	75.75
	ATOM	126	CA	CYS	17	9.582	115.477	35.861	1.00	77.01
	ATOM	127	C	CYS	17	9.032	116.588	36.774	1.00	75.86
	ATOM	128	O	CYS	17	9.603	117.668	36.907	1.00	73.08
	ATOM	129	CB	CYS	17	10.496	114.532	36.638	1.00	80.96
65	ATOM	130	SG	CYS	17	9.859	112.829	36.786	1.00	86.58
	ATOM	131	N	VAL	18	7.895	116.275	37.396	1.00	75.89

	ATOM	132	CA	VAL	18	7.204	117.153	38.296	1.00	77.02
	ATOM	133	C	VAL	18	6.797	116.360	39.498	1.00	78.19
	ATOM	134	O	VAL	18	6.618	115.154	39.447	1.00	77.54
	ATOM	135	CB	VAL	18	5.939	117.816	37.689	1.00	76.68
5	ATOM	136	CG1	VAL	18	4.878	118.021	38.749	1.00	76.28
	ATOM	137	CG2	VAL	18	6.298	119.143	37.052	1.00	76.37
	ATOM	138	N	ALA	19	6.621	117.093	40.600	1.00	80.09
	ATOM	139	CA	ALA	19	6.181	116.535	41.921	1.00	82.16
	ATOM	140	C	ALA	19	4.681	116.512	42.028	1.00	84.28
10	ATOM	141	O	ALA	19	4.070	117.549	41.762	1.00	84.12
	ATOM	142	CB	ALA	19	6.780	117.352	43.067	1.00	80.00
	ATOM	143	N	CYS	20	4.016	115.435	42.343	1.00	87.37
	ATOM	144	CA	CYS	20	2.559	115.616	42.189	1.00	88.33
	ATOM	145	CB	CYS	20	1.853	114.255	42.229	1.00	86.90
15	ATOM	146	SG	CYS	20	1.396	113.592	40.597	1.00	84.56
	ATOM	147	C	CYS	20	1.943	116.641	43.143	1.00	89.64
	ATOM	148	O	CYS	20	0.742	116.890	43.097	1.00	89.77
	ATOM	149	N	GLY	21	2.762	117.223	44.018	1.00	87.37
	ATOM	150	CA	GLY	21	2.224	118.189	44.948	1.00	88.33
20	ATOM	151	C	GLY	21	1.706	119.427	44.214	1.00	89.64
	ATOM	152	O	GLY	21	1.003	120.273	44.761	1.00	89.77
	ATOM	153	N	LEU	22	2.079	119.548	42.934	1.00	87.56
	ATOM	154	CA	LEU	22	1.698	120.688	42.100	1.00	85.33
	ATOM	155	CB	LEU	22	2.712	120.913	40.992	1.00	79.98
25	ATOM	156	CG	LEU	22	3.526	122.194	41.091	1.00	77.91
	ATOM	157	CD1	LEU	22	4.742	122.137	40.187	1.00	78.13
	ATOM	158	CD2	LEU	22	2.656	123.406	40.754	1.00	72.50
	ATOM	159	C	LEU	22	0.320	120.531	41.483	1.00	86.05
	ATOM	160	O	LEU	22	-0.478	121.456	41.559	1.00	88.06
30	ATOM	161	N	LEU	23	0.039	119.361	40.856	1.00	85.11
	ATOM	162	CA	LEU	23	-1.243	119.198	40.162	1.00	83.33
	ATOM	163	C	LEU	23	-2.419	118.778	41.040	1.00	84.21
	ATOM	164	O	LEU	23	-3.542	118.646	40.528	1.00	86.28
	ATOM	165	CB	LEU	23	-1.104	118.191	39.016	1.00	78.93
35	ATOM	166	CG	LEU	23	0.100	118.220	38.087	1.00	73.68
	ATOM	167	CD1	LEU	23	0.216	116.895	37.365	1.00	20.00
	ATOM	168	CD2	LEU	23	0.021	119.370	37.112	1.00	20.00
	ATOM	169	N	ARG	24	-2.218	118.588	42.335	1.00	86.09
	ATOM	170	CA	ARG	24	-3.375	118.141	43.116	1.00	85.39
40	ATOM	171	C	ARG	24	-4.135	119.274	43.764	1.00	84.10
	ATOM	172	O	ARG	24	-5.323	119.128	44.049	1.00	83.24
	ATOM	173	CB	ARG	24	-2.990	117.106	44.135	1.00	80.05
	ATOM	174	CG	ARG	24	-1.599	117.213	44.687	1.00	20.00
	ATOM	175	CD	ARG	24	-1.550	116.457	45.986	1.00	20.00
45	ATOM	176	NE	ARG	24	-0.851	115.207	45.839	1.00	20.00
	ATOM	177	CZ	ARG	24	-0.504	114.518	46.913	1.00	20.00
	ATOM	178	NH1	ARG	24	-0.816	114.966	48.121	1.00	20.00
	ATOM	179	NH2	ARG	24	0.150	113.382	46.778	1.00	20.00
	ATOM	180	N	THR	25	-3.480	120.421	44.031	1.00	84.61
50	ATOM	181	CA	THR	25	-4.187	121.564	44.654	1.00	85.07
	ATOM	182	C	THR	25	-5.030	122.250	43.579	1.00	85.65
	ATOM	183	O	THR	25	-6.086	122.818	43.834	1.00	86.77
	ATOM	184	CB	THR	25	-3.178	122.493	45.356	1.00	20.00
	ATOM	185	OG1	THR	25	-2.315	121.705	46.173	1.00	20.00
55	ATOM	186	CG2	THR	25	-3.907	123.520	46.205	1.00	20.00
	ATOM	187	N	PRO	26	-4.471	122.159	42.329	1.00	84.19
	ATOM	188	CA	PRO	26	-5.192	122.510	41.036	1.00	82.44
	ATOM	189	C	PRO	26	-6.277	121.504	40.607	1.00	84.32
	ATOM	190	O	PRO	26	-7.088	121.734	39.708	1.00	81.24
60	ATOM	191	CB	PRO	26	-4.077	122.602	40.019	1.00	78.00
	ATOM	192	CG	PRO	26	-2.978	123.236	40.801	1.00	20.00
	ATOM	193	CD	PRO	26	-3.263	122.979	42.262	1.00	20.00
	ATOM	194	N	ARG	27	-6.243	120.338	41.297	1.00	87.06
	ATOM	195	CA	ARG	27	-7.019	119.086	41.080	1.00	88.17
65	ATOM	196	C	ARG	27	-6.922	118.739	39.566	1.00	88.45
	ATOM	197	O	ARG	27	-7.790	118.084	38.993	1.00	84.89

	ATOM	198	CB	ARG	27	-8.379	119.094	41.856	0.00	86.84
	ATOM	199	CG	ARG	27	-8.208	118.825	43.405	0.00	85.55
	ATOM	200	CD	ARG	27	-9.093	117.729	44.117	1.00	20.00
	ATOM	201	NE	ARG	27	-8.847	117.550	45.577	1.00	20.00
5	ATOM	202	CZ	ARG	27	-9.432	116.685	46.419	1.00	20.00
	ATOM	203	NH1	ARG	27	-10.345	115.854	45.958	1.00	20.00
	ATOM	204	NH2	ARG	27	-9.103	116.676	47.702	1.00	20.00
	ATOM	205	N	PRO	28	-5.795	119.223	39.005	1.00	89.49
	ATOM	206	CA	PRO	28	-5.388	118.807	37.618	1.00	91.86
10	ATOM	207	C	PRO	28	-5.282	117.244	37.524	1.00	95.75
	ATOM	208	O	PRO	28	-5.137	116.654	36.456	1.00	96.59
	ATOM	209	CB	PRO	28	-4.152	119.672	37.344	0.00	91.23
	ATOM	210	CG	PRO	28	-4.550	120.972	38.003	1.00	20.00
	ATOM	211	CD	PRO	28	-5.556	120.670	39.078	1.00	20.00
15	ATOM	212	N	LYS	29	-5.365	116.616	38.722	1.00	98.41
	ATOM	213	CA	LYS	29	-5.297	115.172	39.038	1.00	96.39
	ATOM	214	C	LYS	29	-4.652	114.160	38.099	1.00	96.88
	ATOM	215	O	LYS	29	-5.212	113.822	37.059	1.00	94.52
	ATOM	216	CB	LYS	29	-6.710	114.693	39.365	0.00	95.29
20	ATOM	217	CG	LYS	29	-7.803	115.599	38.836	0.00	92.34
	ATOM	218	CD	LYS	29	-8.435	115.022	37.585	0.00	89.50
	ATOM	219	CE	LYS	29	-9.851	115.548	37.387	1.00	20.00
	ATOM	220	NZ	LYS	29	-10.805	114.460	37.045	1.00	20.00
	ATOM	221	N	PRO	30	-3.405	113.700	38.473	1.00	97.72
25	ATOM	222	CA	PRO	30	-2.767	112.529	37.778	1.00	98.94
	ATOM	223	C	PRO	30	-3.463	111.181	37.995	1.00	100.71
	ATOM	224	O	PRO	30	-3.613	110.743	39.144	1.00	100.56
	ATOM	225	CB	PRO	30	-1.337	112.546	38.247	0.00	97.33
	ATOM	226	CG	PRO	30	-1.039	113.994	38.324	1.00	20.00
30	ATOM	227	CD	PRO	30	-2.356	114.723	38.449	1.00	20.00
	ATOM	228	N	ALA	31	-3.880	110.511	36.919	1.00	101.17
	ATOM	229	CA	ALA	31	-4.542	109.234	37.046	1.00	99.22
	ATOM	230	C	ALA	31	-3.618	108.090	36.633	1.00	99.01
	ATOM	231	O	ALA	31	-3.863	106.922	36.944	1.00	102.17
35	ATOM	232	CB	ALA	31	-5.815	109.232	36.213	1.00	20.00
	ATOM	233	N	GLY	32	-2.549	108.475	35.961	1.00	95.52
	ATOM	234	CA	GLY	32	-1.544	107.528	35.630	1.00	91.44
	ATOM	235	C	GLY	32	-0.470	107.663	36.696	1.00	88.49
	ATOM	236	O	GLY	32	0.571	108.255	36.455	1.00	86.70
40	ATOM	237	N	ALA	33	-0.725	107.100	37.868	1.00	15.00
	ATOM	238	CA	ALA	33	0.247	107.128	38.944	1.00	15.00
	ATOM	239	C	ALA	33	-0.009	108.145	40.029	1.00	15.00
	ATOM	240	O	ALA	33	0.343	107.933	41.191	1.00	15.00
	ATOM	241	CB	ALA	33	1.640	107.358	38.360	1.00	15.00
45	ATOM	242	N	SER	34	-0.624	109.245	39.684	1.00	15.00
	ATOM	243	CA	SER	34	-0.799	110.309	40.675	1.00	15.00
	ATOM	244	C	SER	34	-1.943	110.091	41.661	1.00	15.00
	ATOM	245	O	SER	34	-1.862	110.483	42.828	1.00	15.00
	ATOM	246	CB	SER	34	-1.001	111.631	39.949	1.00	15.00
50	ATOM	247	OG	SER	34	-0.718	112.722	40.809	1.00	20.00
	ATOM	248	N	SER	35	-3.017	109.472	41.210	1.00	15.00
	ATOM	249	CA	SER	35	-4.197	109.243	42.054	1.00	15.00
	ATOM	250	C	SER	35	-3.758	108.748	43.433	1.00	15.00
	ATOM	251	O	SER	35	-4.296	109.117	44.477	1.00	15.00
55	ATOM	252	CB	SER	35	-5.160	108.240	41.398	1.00	15.00
	ATOM	253	OG	SER	35	-4.769	107.961	40.061	1.00	20.00
	ATOM	254	N	PRO	36	-2.745	107.891	43.359	1.00	15.00
	ATOM	255	CA	PRO	36	-2.102	107.305	44.567	1.00	15.00
	ATOM	256	C	PRO	36	-1.737	108.329	45.608	1.00	15.00
60	ATOM	257	O	PRO	36	-2.129	108.228	46.770	1.00	15.00
	ATOM	258	CB	PRO	36	-1.025	106.412	43.951	1.00	15.00
	ATOM	259	CG	PRO	36	-1.717	105.882	42.720	1.00	15.00
	ATOM	260	CD	PRO	36	-2.753	106.867	42.305	1.00	15.00
	ATOM	261	N	ALA	37	-0.994	109.328	45.169	1.00	15.00
65	ATOM	262	CA	ALA	37	-0.603	110.432	46.018	1.00	15.00
	ATOM	263	C	ALA	37	-1.845	111.170	46.491	1.00	15.00

	ATOM	264	O	ALA	37	-1.902	111.650	47.616	1.00	15.00
	ATOM	265	CB	ALA	37	0.324	111.370	45.269	1.00	15.00
	ATOM	266	N	PRO	38	-2.845	111.293	45.584	1.00	15.00
	ATOM	267	CA	PRO	38	-4.115	112.008	45.935	1.00	15.00
5	ATOM	268	C	PRO	38	-4.777	111.583	47.189	1.00	15.00
	ATOM	269	O	PRO	38	-5.086	112.371	48.071	1.00	15.00
	ATOM	270	CB	PRO	38	-4.971	111.881	44.715	1.00	15.00
	ATOM	271	CG	PRO	38	-3.938	112.068	43.642	1.00	20.00
	ATOM	272	CD	PRO	38	-2.592	111.645	44.189	1.00	20.00
10	ATOM	273	N	ARG	39	-4.977	110.273	47.253	1.00	15.00
	ATOM	274	CA	ARG	39	-5.635	109.668	48.409	1.00	15.00
	ATOM	275	C	ARG	39	-4.649	109.463	49.580	1.00	15.00
	ATOM	276	O	ARG	39	-5.056	108.959	50.631	1.00	15.00
	ATOM	277	CB	ARG	39	-6.276	108.346	48.051	1.00	15.00
15	ATOM	278	CG	ARG	39	-7.190	108.403	46.859	1.00	15.00
	ATOM	279	CD	ARG	39	-6.993	107.148	46.039	1.00	15.00
	ATOM	280	NE	ARG	39	-7.870	107.110	44.888	1.00	20.00
	ATOM	281	CZ	ARG	39	-8.363	105.986	44.378	1.00	20.00
	ATOM	282	NH1	ARG	39	-8.077	104.822	44.944	1.00	20.00
20	ATOM	283	NH2	ARG	39	-9.141	106.030	43.307	1.00	20.00
	ATOM	284	N	THR	40	-3.376	109.815	49.380	1.00	15.00
	ATOM	285	CA	THR	40	-2.270	109.676	50.363	1.00	15.00
	ATOM	286	C	THR	40	-2.233	110.914	51.265	1.00	15.00
	ATOM	287	O	THR	40	-2.185	110.863	52.489	1.00	15.00
25	ATOM	288	CB	THR	40	-0.932	109.558	49.671	1.00	15.00
	ATOM	289	OG1	THR	40	-1.062	108.615	48.595	1.00	20.00
	ATOM	290	CG2	THR	40	0.133	109.086	50.641	1.00	20.00
	ATOM	291	N	ALA	41	-2.267	112.044	50.565	1.00	15.00
	ATOM	292	CA	ALA	41	-2.312	113.313	51.236	1.00	15.00
30	ATOM	293	C	ALA	41	-3.735	113.617	51.697	1.00	15.00
	ATOM	294	O	ALA	41	-3.930	114.488	52.527	1.00	15.00
	ATOM	295	CB	ALA	41	-1.818	114.438	50.332	1.00	15.00
	ATOM	296	N	LEU	42	-4.710	112.932	51.167	1.00	15.00
	ATOM	297	CA	LEU	42	-6.098	113.134	51.563	1.00	15.00
35	ATOM	298	C	LEU	42	-6.357	112.412	52.876	1.00	15.00
	ATOM	299	O	LEU	42	-6.954	112.967	53.804	1.00	15.00
	ATOM	300	CB	LEU	42	-7.052	112.612	50.501	1.00	15.00
	ATOM	301	CG	LEU	42	-7.085	113.368	49.172	1.00	20.00
	ATOM	302	CD1	LEU	42	-8.106	112.746	48.225	1.00	20.00
40	ATOM	303	CD2	LEU	42	-7.387	114.835	49.396	1.00	20.00
	ATOM	304	N	GLN	43	-5.898	111.173	52.968	1.00	15.00
	ATOM	305	CA	GLN	43	-6.011	110.364	54.192	1.00	15.00
	ATOM	306	C	GLN	43	-5.341	111.069	55.347	1.00	15.00
	ATOM	307	O	GLN	43	-5.980	111.430	56.332	1.00	15.00
45	ATOM	308	CB	GLN	43	-5.364	108.976	54.009	1.00	15.00
	ATOM	309	CG	GLN	43	-5.719	108.232	52.743	1.00	20.00
	ATOM	310	CD	GLN	43	-4.929	106.940	52.670	1.00	20.00
	ATOM	311	OE1	GLN	43	-4.062	106.668	53.496	1.00	20.00
	ATOM	312	NE2	GLN	43	-5.091	105.988	51.757	1.00	20.00
50	END					60.588	44.144	42.385	0.00	0.00

This application incorporates by reference in their entirety each of the following provisional applications: U.S. Provisional Application Serial No. 60/345,106, filed October 24, 2001; U.S. Provisional Application Serial No. 60/348,962, filed January 14, 2002; U.S. Provisional Application Serial No. 60/354,966, filed February 7, 2002; and U.S. Provisional
5 Application Serial No. 60/403,364, filed August 13, 2002.

While various embodiments of the present invention have been described in detail, it is apparent that modifications and adaptations of those embodiments will occur to those skilled in the art. It is to be expressly understood, however, that such modifications and adaptations are within the scope of the present invention, as set forth in the following claims.

What is claimed is:

1. A TALL-1 antagonist protein, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one modification in the region connecting β strands D and E that reduces the biological activity of the TALL-1 antagonist as compared to wild-type TALL-1.
2. The TALL-1 antagonist protein of Claim 1, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224.
3. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least two amino acid residues selected from the group consisting of Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224.
4. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least between about 3 and 8 amino acid residues selected from the group consisting of Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224.
5. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least a deletion of the following amino acid residues: Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224.
6. The TALL-1 antagonist protein of Claim 6, wherein the TALL-1 antagonist protein further comprises a substitution of at least one non-natural amino acid residue for said deleted residues.
7. The TALL-1 antagonist protein of Claim 1, wherein the protein has a reduced ability to form a trimer with other TALL-1 monomers.

8. The TALL-1 antagonist protein of Claim 1, wherein the protein, when in a trimer with two other TALL-1 monomers, reduces the ability of the trimer to interact with other TALL-1 trimers.

9. The TALL-1 antagonist protein of Claim 8, wherein each of the two other
5 TALL-1 monomers is selected from the group consisting of: a wild-type TALL-1 monomer and a TALL-1 antagonist protein.

10. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist protein binds to a TALL-1 receptor selected from the group consisting of BCMA, BAFF-R and TACI.

10 11. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said
15 TALL-1 receptor.

12. The TALL-1 antagonist protein of Claim 1, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification in at least one amino acid residue selected from the group consisting of: Tyr163, Tyr206,
20 Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222;

wherein said additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said TALL-1 receptor.

25 13. The TALL-1 antagonist protein of Claim 12, wherein the TALL-1 receptor is selected from the group consisting of BCMA, BAFF-R and TACI.

14. A composition comprising the TALL-1 antagonist protein of Claim 1.

15. A TALL-1 antagonist protein, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of
30 positions 134 to 285 of SEQ ID NO:2, by at least one modification that reduces interaction between a first trimer and a second trimer, wherein said first trimer comprises:

- a. a monomer of said TALL-1 antagonist protein; and
- b. two monomers selected from the group consisting of: wild-type TALL-1 monomers, said TALL-1 antagonist protein monomers, and mixtures thereof;

5 and wherein said second trimer comprises monomers selected from the group consisting of wild-type TALL-1 monomers, said TALL-1 antagonist protein monomers, and mixtures thereof.

16. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence
10 consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue located in a region of TALL-1 selected from the group consisting of β strand C, β strand F, and the region connecting β strand D to β strand E.

17. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence
15 consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Ile150, Leu169, Phe172, Tyr192, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Ile250, Lys252, and Glu254.

18. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises
20 an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, and Leu224.

19. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises
25 an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Tyr192, Lys252, Glu254, His218, Lys216, Glu223, Leu224, Val227, Leu229, Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172.

30 20. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence

consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Tyr192, Lys252, Glu254, and His218.

21. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence
5 consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Lys216, Glu223, Leu224, Val227, and Leu229.

22. The TALL-1 antagonist protein of Claim 15, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence
10 consisting of positions 134 to 285 of SEQ ID NO:2, by a modification in at least one amino acid residue selected from the group consisting of: Val219, Ile150, Leu169, Phe220, Tyr192, Ile250 and Phe172.

23. The TALL-1 antagonist protein of Claim 15, wherein the TALL-1 antagonist protein binds to a TALL-1 receptor selected from the group consisting of BCMA, BAFF-R
15 and TACI.

24. The TALL-1 antagonist protein of Claim 15, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and
20 a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said TALL-1 receptor.

25. The TALL-1 antagonist protein of Claim 15, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification
25 in at least one amino acid residue selected from the group consisting of: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222;

wherein said additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity
30 between wild-type TALL-1 and said TALL-1 receptor.

26. The TALL-1 antagonist protein of Claim 25, wherein the TALL-1 receptor is selected from the group consisting of BCMA, BAFF-R and TACI.

27. The TALL-1 antagonist protein of Claim 15, wherein the protein has a reduced ability to form a trimer with other TALL-1 monomers.

5 28. A composition comprising the TALL-1 antagonist protein of Claim 15.

29. A TALL-1 antagonist protein, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue selected from the group consisting of: Phe194, Tyr196, Tyr246,
10 Leu282, Gln144 and Leu285.

30. The TALL-1 antagonist protein of Claim 29, wherein the protein has a reduced ability to form a trimer with other TALL-1 monomers.

31. The TALL-1 antagonist protein of Claim 29, wherein the TALL-1 antagonist protein binds to a TALL-1 receptor selected from the group consisting of BCMA, BAFF-R
15 and TACI.

32. The TALL-1 antagonist protein of Claim 29, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by at least one additional modification that increases the binding affinity between the TALL-1 antagonist protein and
20 a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said TALL-1 receptor.

33. The TALL-1 antagonist protein of Claim 29, wherein the TALL-1 antagonist comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by an additional modification
25 in at least one amino acid residue selected from the group consisting of: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222;

wherein the additional modification increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between
30 wild-type TALL-1 and said TALL-1 receptor.

34. The TALL-1 antagonist protein of Claim 33, wherein the TALL-1 receptor is selected from the group consisting of BCMA, BAFF-R and TACI.

35. A composition comprising the TALL-1 antagonist protein of Claim 29.

36. A TALL-1 antagonist protein, wherein the TALL-1 antagonist protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue that reduces the biological activity of the antagonist protein as compared to a wild-type TALL-1, wherein said amino acid residue is selected from the group consisting of: Gln144, Ile150, Leu169, Phe172, Tyr192, Phe194, Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285; and

wherein the amino acid sequence of the TALL-1 antagonist further differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification of at least one amino acid residue that increases the binding affinity between the TALL-1 antagonist protein and a TALL-1 receptor, as compared to the binding affinity between wild-type TALL-1 and said TALL-1 receptor, wherein said amino acid residue is selected from the group consisting of: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222.

37. A composition comprising the TALL-1 antagonist protein of Claim 36.

38. A TALL-1 antagonist protein, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from the group consisting of: Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222;

wherein the TALL-1 antagonist protein has reduced binding to a receptor for TALL-1 as compared to wild-type TALL-1.

39. The TALL-1 antagonist protein of Claim 38, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from the group consisting of: Tyr163, Leu211, Ile233, Pro264, and Leu200.

40. The TALL-1 antagonist protein of Claim 38, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from the group consisting of: Tyr206 and Leu240.

5 41. The TALL-1 antagonist protein of Claim 38, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from the group consisting of: Arg265, Glu266 and Glu238.

10 42. The TALL-1 antagonist protein of Claim 38, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:2, or from an amino acid sequence consisting of positions 134 to 285 of SEQ ID NO:2, by a modification to at least one amino acid residue selected from the group consisting of: Asp222, Asp 273 and Asp275.

43. The TALL-1 antagonist protein of Claim 38, wherein the TALL-1 receptor is selected from the group consisting of BCMA, BAFF-R, and TACI.

15 44. The TALL-1 antagonist protein of Claim 38, wherein the TALL-1 antagonist protein has reduced ability to bind to at least two of BCMA, BAFF-R and TACI.

45. The TALL-1 antagonist protein of Claim 38, wherein the TALL-1 antagonist protein has reduced ability to bind to each of BCMA, BAFF-R and TACI.

46. A composition comprising the TALL-1 antagonist protein of Claim 38.

20 47. An APRIL agonist protein, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:4 by at least one modification that increases the binding affinity between the APRIL agonist protein and an APRIL receptor, as compared to the binding affinity between wild-type APRIL and said APRIL receptor.

25 48. The APRIL agonist protein of Claim 47, wherein said protein comprises an amino acid sequence that differs from SEQ ID NO:4 by a modification in at least one amino acid residue selected from the group consisting of: Val133, Thr177, Val181, Ile197, Pro230, Leu58, Tyr96, Phe176, Arg206, and Arg265;

wherein the modification increases the binding affinity between the APRIL agonist protein and an APRIL receptor, as compared to the binding affinity between wild-type
30 APRIL and said APRIL receptor.

49. The APRIL agonist protein of Claim 47, wherein said APRIL receptor is selected from the group consisting of BCMA and TACI.

50. The APRIL agonist protein of Claim 47, wherein said at least one modification results in binding of said APRIL to BAFF-R.

5 51. A method to inhibit TALL-1 biological activity in a mammal, comprising administering to said mammal the protein of any one of Claims 1, 15, 29, 36 or 47.

52. The method of Claim 51, wherein said protein is a competitive inhibitor of wild-type TALL-1 for binding to a TALL-1 receptor.

53. The method of Claim 51, wherein the mammal has, or is at risk of developing,
10 a disease or condition associated with hyperactive B cell development or B cell hyperproliferation.

54. The method of Claim 51, wherein the mammal has, or is at risk of developing, a disease or condition characterized by increased numbers of mature B-lymphocytes, splenomegaly, anti-DNA antibodies, proteinuria, or glomerulonephritis.

15 55. The method of Claim 54, wherein said disease is systemic lupus erythematosus.

56. A recombinant nucleic acid molecule comprising a nucleic acid sequence encoding the amino acid sequence of any one of Claims 1, 15, 29, 36, or 47 operatively linked to a transcription control sequence.

20 57. A method to inhibit TALL-1 biological activity in a mammal, comprising administering to said mammal the recombinant nucleic acid molecule of Claim 56, wherein said protein is expressed by a host cell in said mammal.

58. The method of Claim 57, wherein said protein associates with wild-type TALL-1 monomers expressed by said cell to produce TALL-1 trimers containing said protein
25 with reduced TALL-1 biological activity, as compared to a trimer of wild-type TALL-1 monomers.

59. A recombinant nucleic acid molecule comprising a nucleic acid sequence encoding the amino acid sequence of Claim 38, operatively linked to a transcription control sequence.

60. A method to inhibit TALL-1 biological activity in a mammal, comprising administering to said mammal the recombinant nucleic acid molecule of Claim 59, wherein said protein is expressed by a host cell in said mammal.

61. The method of Claim 60, wherein said protein associates with wild-type
5 TALL-1 monomers expressed by said cell to produce TALL-1 trimers containing said protein with reduced ability to bind to a TALL-1 receptor, as compared to a trimer of wild-type TALL-1 monomers.

62. A BCMA antagonist, wherein said receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:6 by a modification in at least one amino acid
10 residue selected from the group consisting of: Tyr13, Asp15, Leu17, Leu18, His19, Ile22, Leu26, Arg27, and Pro34;

wherein said BCMA antagonist has an increased binding affinity for TALL-1 as compared to wild-type BCMA.

63. The BCMA antagonist of Claim 62, wherein said amino acid residue is
15 selected from the group consisting of Leu17 and Leu18.

64. The BCMA antagonist of Claim 62, wherein said amino acid residue is selected from the group consisting of Ile22 and Leu26.

65. The BCMA antagonist of Claim 62, wherein said amino acid residue is selected from the group consisting of Asp15, Arg27 and Tyr13.

20 66. The BCMA antagonist of Claim 62, wherein said amino acid residue is His19.

67. The BCMA antagonist of Claim 62, wherein said amino acid residue is selected from the group consisting of Tyr13, Leu17, Leu18 and Ile22.

68. The BCMA antagonist of Claim 67, wherein said amino acid residue is substituted with an amino acid residue selected from the group consisting of: Ile, Met, Phe
25 or Tyr.

69. The BCMA antagonist of Claim 62, wherein said BCMA antagonist is a soluble protein.

70. A BAFF-R antagonist, wherein said receptor antagonist comprises an amino acid sequence that differs from SEQ ID NO:8 by a modification in at least one amino acid
30 residue selected from the group consisting of: Asp26, Leu28, Val29, Arg30, Val33, Leu37, Leu38, and Arg42, and Pro45;

wherein said BAFF-R antagonist has an increased binding affinity for TALL-1 as compared to wild-type BAFF-R.

71. The BAFF-R antagonist of Claim 70, wherein said amino acid residue is selected from the group consisting of Leu28 and Val29.

5 72. The BAFF-R antagonist of Claim 70, wherein said amino acid residue is selected from the group consisting of Val33, Leu37, Leu38 and Pro45.

73. The BAFF-R antagonist of Claim 70, wherein said amino acid residue is selected from the group consisting of Asp26 and Arg 42.

74. The BAFF-R antagonist of Claim 70, wherein said amino acid residue is
10 selected from the group consisting of Arg30.

75. The BAFF-R antagonist of Claim 70, wherein said amino acid residue is selected from the group consisting of Leu28, Val29 and Val33.

76. The BAFF-R antagonist of Claim 75, wherein said amino acid residue is substituted with an amino acid residue selected from the group consisting of: Ile, Met, Phe
15 or Tyr.

77. The BAFF-R antagonist of Claim 70, wherein said BAFF-R antagonist is a soluble protein.

78. A method to inhibit TALL-1 receptor biological activity in a mammal, comprising administering to said mammal the antagonist of any one of Claims 62 or 70.

20 79. The method of Claim 78, wherein said antagonist is a competitive inhibitor of a wild-type TALL-1 receptor for binding to TALL-1.

80. A method to inhibit the biological activity of TALL-1, comprising administering to a cell that expresses TALL-1 a recombinant nucleic acid molecule comprising a nucleic acid sequence encoding APRIL, or a biologically active fragment
25 thereof.

81. An isolated BAFF-R antagonist, wherein said BAFF-R antagonist consists essentially of the amino acid sequence represented by SEQ ID NO:9, or homologues thereof with substantially the same biological activity.

82. A method to identify a compound that is a competitive inhibitor of TALL-1
30 binding to its receptor, comprising:

a. contacting a TALL-1 receptor or a TALL-1 binding fragment thereof with a homologue of a TALL-1 protein, wherein said homologue comprises an amino acid sequence with a modification in at least one amino acid residue selected from the group consisting of Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, and Glu238; and

b. detecting whether said homologue binds to said TALL-1 receptor or fragment thereof;

wherein homologues that bind to said TALL-1 receptor or fragment thereof potential competitive inhibitors for binding of wild-type TALL-1 to its receptor.

83. The method of Claim 82, further comprising a step (c) of detecting whether homologues that bind to said TALL-1 receptor or fragment thereof in (b) have a TALL-1 biological activity selected from the group consisting of: an ability to activate signal transduction in said TALL-1 receptor, an ability to form a trimer with two other TALL-1 monomers, an ability to form a trimer with TALL-1 two other TALL-1 monomers that is capable of interacting with other TALL-1 trimers;

wherein homologues that have a decreased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 antagonists, and wherein homologues that have an increased TALL-1 biological activity as compared to wild-type TALL-1 are identified as TALL-1 agonists.

84. The method of Claim 83, wherein step (b) further comprises comparing the binding affinity said homologue to said TALL-1 receptor or fragment of thereof to the binding affinity of wild-type TALL-1 and said TALL-1 receptor;

and wherein said method further comprises step (d) of selecting homologues which have an increased binding affinity to said TALL-1 receptor or fragment of and a decreased TALL-1 biological activity.

85. A method of structure-based identification of compounds which potentially bind to TALL-1, comprising:

a. obtaining atomic coordinates that define the three dimensional structure of TALL-1, said atomic coordinates being selected from the group consisting of:

i. atomic coordinates determined by X-ray diffraction of a crystalline TALL-1;

ii. atomic coordinates selected from the group consisting of:

(1) atomic coordinates represented in any one of Tables 2-12;

(2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by said atomic coordinates of (1); and

(3) atomic coordinates in any one of Tables 2-12 defining a portion of said TALL-1, wherein the portion of said TALL-1 comprises sufficient structural information to perform step (b); and

iii. atomic coordinates defining the three dimensional structure of TALL-1 molecules arranged in a crystalline manner in a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217\text{\AA}$; and

b. selecting candidate compounds for binding to said TALL-1 by performing structure based drug design with said structure of (a), wherein said step of selecting is performed in conjunction with computer modeling.

86. The method of Claim 85, wherein said method further comprises:

c. selecting candidate compounds of (b) that inhibit the biological activity of TALL-1.

87. The method of Claim 86, wherein said step (c) of selecting comprises:

i. contacting said candidate compound identified in step (b) with TALL-1; and

ii. measuring the biological activity of said TALL-1, as compared to in the absence of said candidate compound.

88. The method of Claim 85, wherein said method further comprises:

c. selecting candidate compounds of (b) that inhibit the binding of TALL-1 to a TALL-1 receptor.

89. The method of Claim 88, wherein said step (c) of selecting comprises:

i. contacting said candidate compound identified in step (b) with said TALL-1 or a fragment thereof and a TALL-1 receptor or TALL-1 receptor binding fragment thereof under conditions in which a TALL-1-TALL-1 receptor complex can form in the absence of said candidate compound; and

ii. measuring the binding of said TALL-1 or fragment thereof to bind to said TALL-1 receptor or fragment thereof, wherein a candidate inhibitor compound is selected when there is a decrease in the binding of the TALL-1 or fragment thereof to the TALL-1 receptor or fragment thereof, as compared to in the absence of said candidate inhibitor compound.

90. The method of Claim 89, wherein said TALL-1 receptor is selected from the group consisting of BCMA, BAFF-R and TACI.

91. The method of Claim 85, wherein said step of selecting comprises identifying candidate compounds for binding to a receptor binding site of said TALL-1 protein, said receptor binding site comprising an amino acid residue selected from the group consisting of Tyr163, Tyr206, Leu211, Arg231, Ile233, Pro264, Arg265, Glu266, Leu200, Leu240, Asp273, Asp275, Glu238 and Asp222.

92. The method of Claim 85, wherein said step of selecting comprises identifying candidate compounds for binding to said TALL-1 such that trimer-trimer interactions between trimers of TALL-1 monomers is inhibited.

93. The method of Claim 92, wherein said step of selecting comprises identifying candidate compounds for binding to TALL-1 at a site including an amino acid residue selected from the group consisting of: Gln144, Ile150, Leu169, Phe172, Tyr192, Phe194, Tyr196, Lys216, Val217, His218, Val219, Phe220, Glu221, Asp222, Glu223, Leu224, Val227, Leu229, Tyr246, Ile250, Lys252, Glu254, Leu282, and Leu285.

94. A therapeutic composition comprising a compound that inhibits the biological activity of TALL-1, said compound being identified by the method of Claim 85.

95. A method to treat a disease or condition that can be regulated by modifying the biological activity of TALL-1, comprising administering to a mammal with such a disease or condition the therapeutic composition of Claim 94.

96. A method to construct a three dimensional model of TALL-1 protein or
5 homologue thereof, comprising:

a. obtaining atomic coordinates that define the three dimensional structure of TALL-1, said atomic coordinates being selected from the group consisting of:

- 10 i. atomic coordinates determined by X-ray diffraction of a crystalline TALL-1;
 - ii. atomic coordinates selected from the group consisting of:
 - (1) atomic coordinates represented in any one of Tables 2-12;
 - 15 (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by said atomic coordinates of (1); and
 - 20 (3) atomic coordinates in any one of Tables 2-12 defining a portion of said TALL-1, wherein the portion of said TALL-1 comprises sufficient structural information to perform step (b); and
 - 25 iii. atomic coordinates defining the three dimensional structure of TALL-1 molecules arranged in a crystalline manner in a space group P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217 Å; and
- a. performing computer modeling with said atomic coordinates of (a) and to construct a model of a three dimensional structure of a TALL-1 or homologue
30 thereof.

97. A method of structure-based identification of compounds which potentially bind to a TALL-1 receptor selected from the group consisting of BCMA and BAFF-R, comprising:

5 a. obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R, said atomic coordinates being selected from the group consisting of:

i. atomic coordinates determined by X-ray diffraction of a crystalline BCMA or crystalline BAFF-R;

10 ii. atomic coordinates selected from the group consisting of:

(1) atomic coordinates represented in any one of Tables 13-33;

15 (2) atomic coordinates that define a three dimensional structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by said atomic coordinates of (1); and

20 (3) atomic coordinates in any one of Tables 13-22 defining a portion of said BCMA, wherein the portion of said BCMA comprises sufficient structural information to perform step (b);

25 (4) atomic coordinates in any one of Tables 14-33 defining a portion of said BAFF-R, wherein the portion of said BAFF-R comprises sufficient structural information to perform step (b); and

iii. atomic coordinates defining the three dimensional structure of BCMA molecules or BAFF-R molecules arranged in a crystalline manner in a space group P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217; and

b. selecting candidate compounds for binding to said BCMA or BAFF-R by performing structure based drug design with said structure of (a), wherein said step of selecting is performed in conjunction with computer modeling.

98. A method to construct a three dimensional model of BCMA, BAFF-R, TACI,
5 or a homologue thereof, comprising:

a. obtaining atomic coordinates that define the three dimensional structure of BCMA or BAFF-R, said atomic coordinates being selected from the group consisting of:

i. atomic coordinates determined by X-ray diffraction of a
10 crystalline BCMA or crystalline BAFF-R;

ii. atomic coordinates selected from the group consisting of:

(1) atomic coordinates represented in any one of Tables 13-33;

(2) atomic coordinates that define a three dimensional
15 structure having an average root-mean-square deviation (RMSD) of equal to or less than about 1.7Å over the backbone atoms in secondary structure elements of at least 50% of the residues in a three dimensional structure represented by said atomic coordinates of (1); and

(3) atomic coordinates in any one of Tables 13-22
20 defining a portion of said BCMA, wherein the portion of said BCMA comprises sufficient structural information to perform step (b);

(4) atomic coordinates in any one of Tables 14-33
25 defining a portion of said BAFF-R, wherein the portion of said BAFF-R comprises sufficient structural information to perform step (b); and

iii. atomic coordinates defining the three dimensional structure of BCMA molecules or BAFF-R molecules arranged in a crystalline manner in
30 a space group $P6_322$ so as to form a unit cell having approximate dimensions of $a=b=234\text{\AA}$, $c=217$; and

b. performing computer modeling with said atomic coordinates of (a) and an amino acid sequence corresponding to BCMA, BAFF-R or TACI to construct a model of a three dimensional structure of said BCMA, BAFF-R or TACI, or homologue thereof.

- 5 99. A crystal comprising a TALL-1 protein, wherein the crystal effectively diffracts X-rays for the determination of the atomic coordinates of the TALL-1 protein to a resolution of greater than 3.0 Å, and P6₃22 so as to form a unit cell having approximate dimensions of a=b=234Å, c=217.

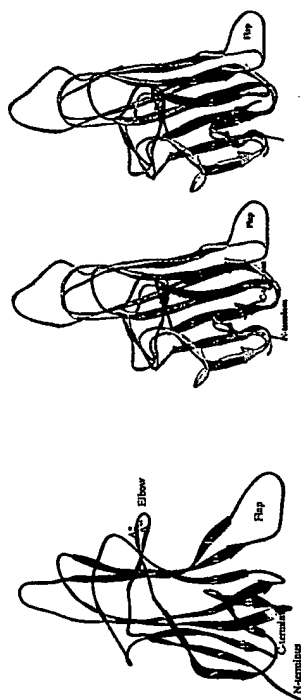


FIG. 1A

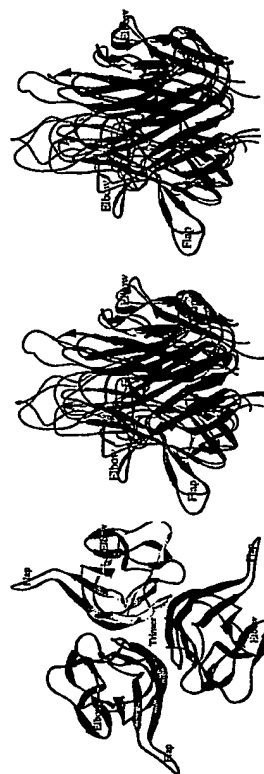


FIG. 1B

FIG. 1C

FIG. 1D

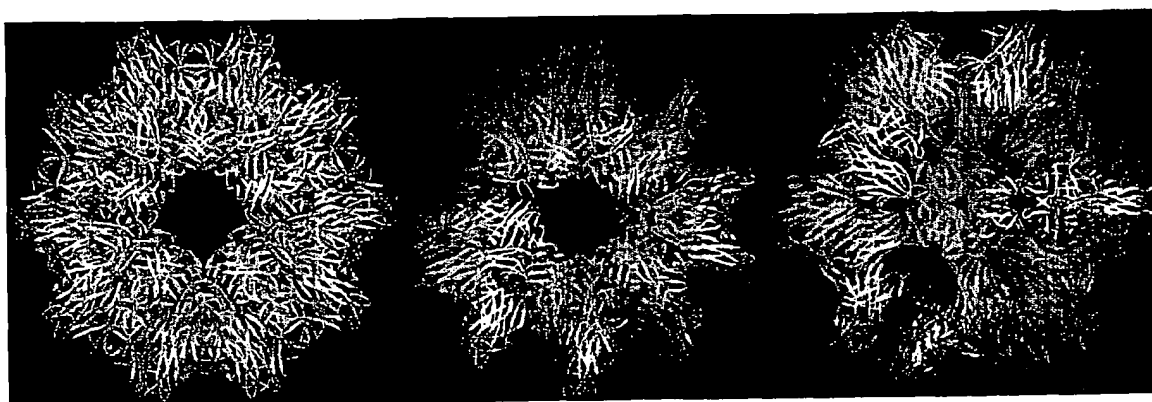


FIG. 2A

FIG. 2B

FIG. 2C

FIG. 3B

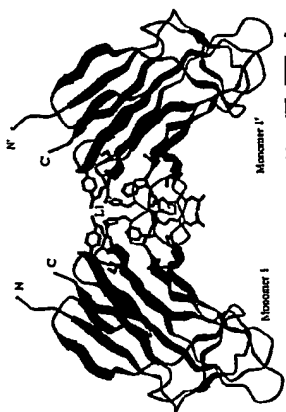


FIG. 3D

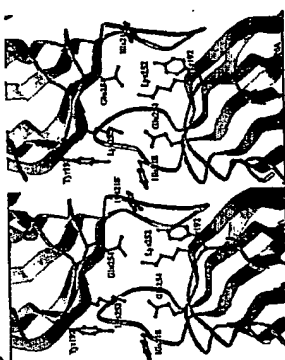


FIG. 3F

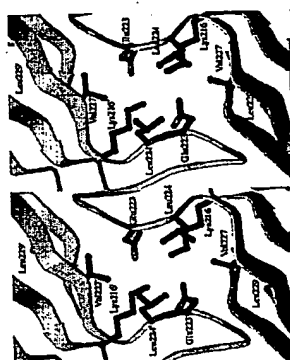


FIG. 3H



FIG. 3A



FIG. 3C



FIG. 3E



FIG. 3G



FIG. 4C

FIG. 4B

FIG. 4A

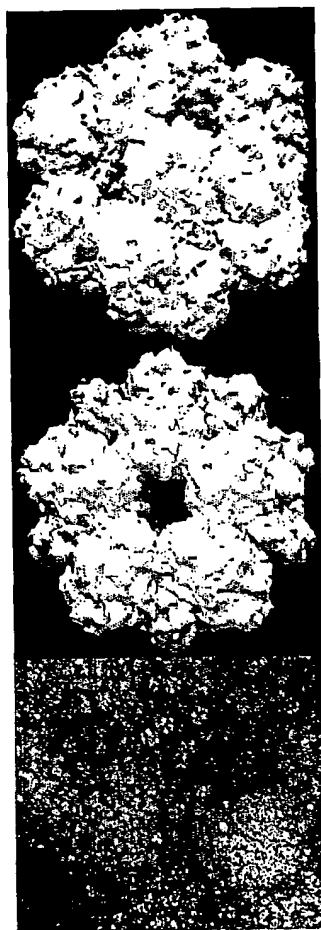




FIG. 5A



FIG. 5B



FIG. 6A

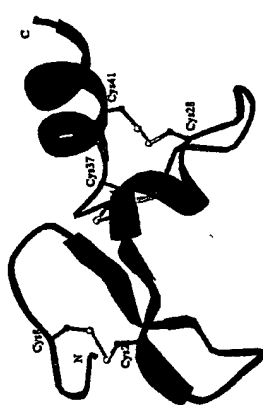


FIG. 6B

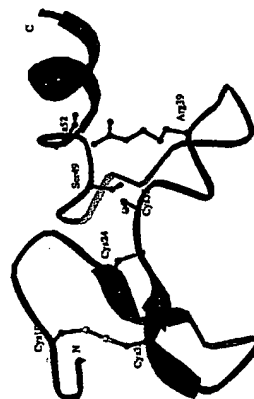


FIG. 6C

FIG. 7A

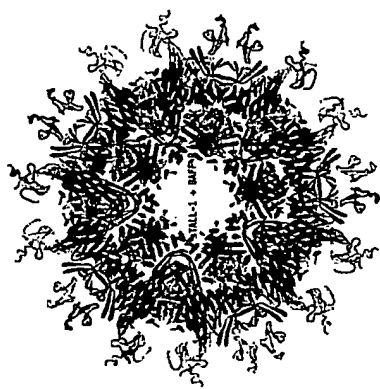


FIG. 7B



FIG. 7C

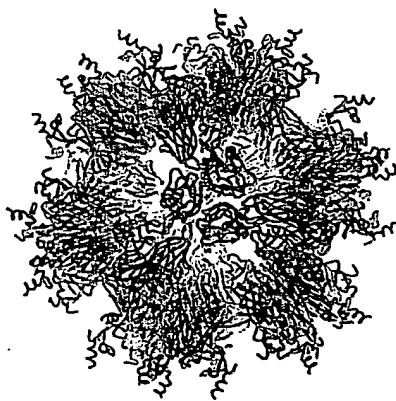


FIG. 7D



FIG. 8A



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 TACII 71 CQGGQTHLLRCSSTQCHK CCVVF 104
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FIG. 8B

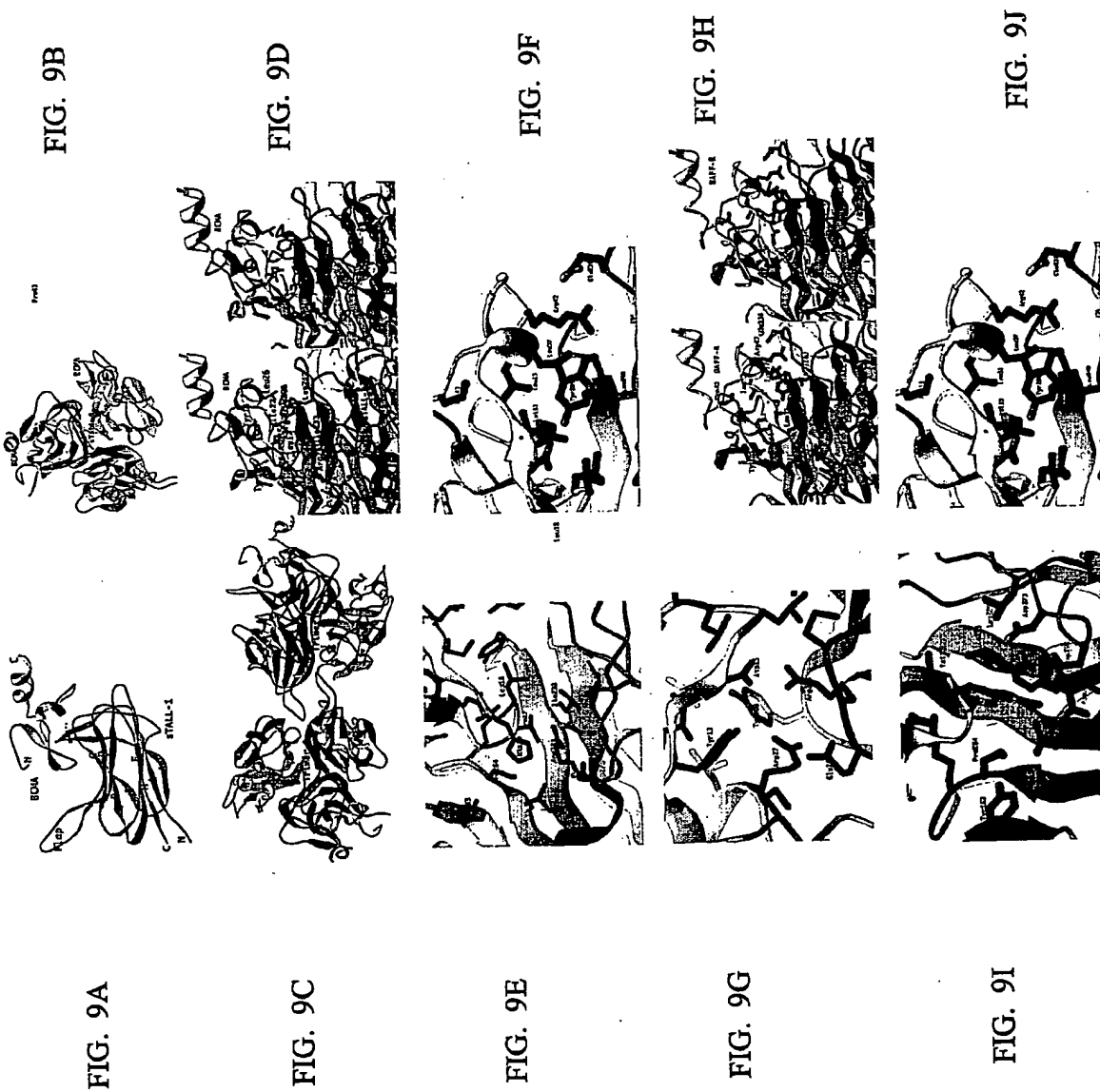




FIG. 10B

FIG. 10A

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<110> Zhang, Gongyi
 Shu, Hong-Bing
 Liu, Yingfang
 Xu, Liangguo

<120> Three-Dimensional Structures of TALL-1 and its Cognate Receptors and Modified Proteins and Methods Related Thereto

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Lys Ile Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn
 115 120 125

Ser Arg Asn Lys Arg Ala Val Gln Gly Pro Glu Glu Thr Val Thr Gln
 130 135 140

Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys
 145 150 155 160

Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser
 165 170 175

Ala Leu Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr
 180 185 190

Phe Phe Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met
 195 200 205

Gly His Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu
 210 215 220

Ser Leu Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu

225					230					235				240
Pro	Asn	Asn	Ser	Cys	Tyr	Ser	Ala	Gly	Ile	Ala	Lys	Leu	Glu	Glu
				245					250					255
Asp	Glu	Leu	Gln	Leu	Ala	Ile	Pro	Arg	Glu	Asn	Ala	Gln	Ile	Ser
			260					265					270	Leu
Asp	Gly	Asp	Val	Thr	Phe	Phe	Gly	Ala	Leu	Lys	Leu	Leu		
		275					280					285		

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<210> 3
<211> 753
<212> DNA
<213> Homo sapiens
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<220>
<221> CDS
<222> (1) .. (753)
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Met	Pro	Ala	Ser	Ser	Pro	Phe	Leu	Leu	Ala	Pro	Lys	Gly	Pro	Pro	Gly	
1				5					10					15		
aac	atg	ggg	ggc	cca	gtc	aga	gag	ccg	gca	ctc	tca	gtt	gcc	ctc	tgg	
Asn	Met	Gly	Gly	Pro	Val	Arg	Glu	Pro	Ala	Leu	Ser	Val	Ala	Leu	Trp	
			20					25					30			
ttg	agt	tgg	ggg	gca	gct	ctg	ggg	gcc	gtg	gct	tgt	gcc	atg	gct	ctg	
Leu	Ser	Trp	Gly	Ala	Ala	Leu	Gly	Ala	Val	Ala	Cys	Ala	Met	Ala	Leu	
		35					40					45				
ctg	acc	caa	caa	aca	gag	ctg	cag	agc	ctc	agg	aga	gag	gtg	agc	cgg	
Leu	Thr	Gln	Gln	Thr	Glu	Leu	Gln	Ser	Leu	Arg	Arg	Glu	Val	Ser	Arg	
	50					55					60					
ctg	cag	ggg	aca	gga	ggc	ccc	tcc	cag	aat	ggg	gaa	ggg	tat	ccc	tgg	
Leu	Gln	Gly	Thr	Gly	Gly	Pro	Ser	Gln	Asn	Gly	Glu	Gly	Tyr	Pro	Trp	
65					70					75					80	
cag	agt	ctc	ccg	gag	cag	agt	tcc	gat	gcc	ctg	gaa	gcc	tgg	gag	aat	
Gln	Ser	Leu	Pro	Glu	Gln	Ser	Ser	Asp	Ala	Leu	Glu	Ala	Trp	Glu	Asn	
				85					90					95		
ggg	gag	aga	tcc	cgg	aaa	agg	aga	gca	gtg	ctc	acc	caa	aaa	cag	aag	
Gly	Glu	Arg	Ser	Arg	Lys	Arg	Arg	Ala	Val	Leu	Thr	Gln	Lys	Gln	Lys	
			100					105					110			
aag	cag	cac	tct	gtc	ctg	cac	ctg	gtt	ccc	att	aac	gcc	acc	tcc	aag	
Lys	Gln	His	Ser	Val	Leu	His	Leu	Val	Pro	Ile	Asn	Ala	Thr	Ser	Lys	
		115					120					125				

gat gac tcc gat gtg aca gag gtg atg tgg caa cca gct ctt agg cgt 432
 Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg Arg
 130 135 140

ggg aga ggc cta cag gcc caa gga tat ggt gtc cga atc cag gat gct 480
 Gly Arg Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp Ala
 145 150 155 160

gga gtt tat ctg ctg tat agc cag gtc ctg ttt caa gac gtg act ttc 528
 Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr Phe
 165 170 175

acc atg ggt cag gtg gtg tct cga gaa ggc caa gga agg cag gag act 576
 Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr
 180 185 190

cta ttc cga tgt ata aga agt atg ccc tcc cac ccg gac cgg gcc tac 624
 Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr
 195 200 205

aac agc tgc tat agc gca ggt gtc ttc cat tta cac caa ggg gat att 672
 Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile
 210 215 220

ctg agt gtc ata att ccc cgg gca agg gcg aaa ctt aac ctc tct cca 720
 Leu Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser Pro
 225 230 235 240

cat gga acc ttc ctg ggg ttt gtg aaa ctg tga 753
 His Gly Thr Phe Leu Gly Phe Val Lys Leu
 245 250

<210> 4
 <211> 250
 <212> PRT
 <213> Homo sapiens

<400> 4

Met Pro Ala Ser Ser Pro Phe Leu Leu Ala Pro Lys Gly Pro Pro Gly
 1 5 10 15

Asn Met Gly Gly Pro Val Arg Glu Pro Ala Leu Ser Val Ala Leu Trp
 20 25 30

Leu Ser Trp Gly Ala Ala Leu Gly Ala Val Ala Cys Ala Met Ala Leu
 35 40 45

Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg Glu Val Ser Arg
 50 55 60

Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly Glu Gly Tyr Pro Trp
 65 70 75 80

Gln Ser Leu Pro Glu Gln Ser Ser Asp Ala Leu Glu Ala Trp Glu Asn
85 90 95

Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln Lys
100 105 110

Lys Gln His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser Lys
115 120 125

Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg Arg
130 135 140

Gly Arg Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp Ala
145 150 155 160

Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr Phe
165 170 175

Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr
180 185 190

Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr
195 200 205

Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile
210 215 220

Leu Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser Pro
225 230 235 240

His Gly Thr Phe Leu Gly Phe Val Lys Leu
245 250

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<211> 995
<212> DNA
<213> Homo sapiens

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<222> (219)..(773)
<223>

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acacagacag cccccgtaag aacccacgaa gcaggcgaag ttcattgttc tcaacattct	120
agctgctctt gctgcatttg ctctggaatt cttgtagaga tattacttgt ccttccaggc	180
tgttctttct gtagctccct tgttttcttt ttgtgatc atg ttg cag atg gct ggg Met Leu Gln Met Ala Gly 1 5	236
cag tgc tcc caa aat gaa tat ttt gac agt ttg ttg cat gct tgc ata Gln Cys Ser Gln Asn Glu Tyr Phe Asp Ser Leu Leu His Ala Cys Ile 10 15 20	284
cct tgt caa ctt cga tgt tct tct aat act cct cct cta aca tgt cag Pro Cys Gln Leu Arg Cys Ser Ser Asn Thr Pro Pro Leu Thr Cys Gln 25 30 35	332
cgt tat tgt aat gca agt gtg acc aat tca gtg aaa gga acg aat gcg Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser Val Lys Gly Thr Asn Ala 40 45 50	380
att ctc tgg acc tgt ttg gga ctg agc tta ata att tct ttg gca gtt Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu Ile Ile Ser Leu Ala Val 55 60 65 70	428
ttc gtg cta atg ttt ttg cta agg aag ata agc tct gaa cca tta aag Phe Val Leu Met Phe Leu Leu Arg Lys Ile Ser Ser Glu Pro Leu Lys 75 80 85	476
gac gag ttt aaa aac aca gga tca ggt ctc ctg ggc atg gct aac att Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu Leu Gly Met Ala Asn Ile 90 95 100	524
gac ctg gaa aag agc agg act ggt gat gaa att att ctt ccg aga ggc Asp Leu Glu Lys Ser Arg Thr Gly Asp Glu Ile Ile Leu Pro Arg Gly 105 110 115	572
ctc gag tac acg gtg gaa gaa tgc acc tgt gaa gac tgc atc aag agc Leu Glu Tyr Thr Val Glu Glu Cys Thr Cys Glu Asp Cys Ile Lys Ser 120 125 130	620
aaa ccg aag gtc gac tct gac cat tgc ttt cca ctc cca gct atg gag Lys Pro Lys Val Asp Ser Asp His Cys Phe Pro Leu Pro Ala Met Glu 135 140 145 150	668
gaa ggc gca acc att ctt gtc acc acg aaa acg aat gac tat tgc aag Glu Gly Ala Thr Ile Leu Val Thr Thr Lys Thr Asn Asp Tyr Cys Lys 155 160 165	716
agc ctg cca gct gct ttg agt gct acg gag ata gag aaa tca att tct Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu Ile Glu Lys Ser Ile Ser 170 175 180	764
gct agg taa ttaaccattt cgactcgagc agtgccactt taaaaatctt Ala Arg	813
ttgtcagaat agatgatgtg tcagatctct ttaggatgac tgtatttttc agttgccgat	873
acagcttttt gtccctctaac tgtggaaact ctttatgtta gatatatctt tctagggtac	933

tggtgggagc ttaatggtag aaacttcctt gggttcatga ttaaagtctt tttttttcct 993
ga 995

<210> 6
<211> 184
<212> PRT
<213> Homo sapiens

<400> 6

Met Leu Gln Met Ala Gly Gln Cys Ser Gln Asn Glu Tyr Phe Asp Ser
1 5 10 15

Leu Leu His Ala Cys Ile Pro Cys Gln Leu Arg Cys Ser Ser Asn Thr
20 25 30

Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser
35 40 45

Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu
50 55 60

Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe Leu Leu Arg Lys Ile
65 70 75 80

Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu
85 90 95

Leu Gly Met Ala Asn Ile Asp Leu Glu Lys Ser Arg Thr Gly Asp Glu
100 105 110

Ile Ile Leu Pro Arg Gly Leu Glu Tyr Thr Val Glu Glu Cys Thr Cys
115 120 125

Glu Asp Cys Ile Lys Ser Lys Pro Lys Val Asp Ser Asp His Cys Phe
130 135 140

Pro Leu Pro Ala Met Glu Glu Gly Ala Thr Ile Leu Val Thr Thr Lys
145 150 155 160

Thr Asn Asp Tyr Cys Lys Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu
165 170 175

Ile Glu Lys Ser Ile Ser Ala Arg
180

<210> 7
 <211> 899
 <212> DNA
 <213> Homo sapiens

<220>
 <221> CDS
 <222> (6)..(560)
 <223>

<400> 7
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 Met Arg Arg Gly Pro Arg Ser Leu Arg Gly Arg Asp Ala Pro Ala
 1 5 10 15

ccc acg ccc tgc gtc ccg gcc gag tgc ttc gac ctg ctg gtc cgc cac 98
 Pro Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg His
 20 25 30

tgc gtg gcc tgc ggg ctc ctg cgc acg ccg cgg ccg aaa ccg gcc ggg 146
 Cys Val Ala Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro Ala Gly
 35 40 45

gcc agc agc cct gcg ccc agg acg gcg ctg cag ccg cag gag tcg gtg 194
 Ala Ser Ser Pro Ala Pro Arg Thr Ala Leu Gln Pro Gln Glu Ser Val
 50 55 60

ggc gcg ggg gcc ggc gag gcg gcg ctg ccc ctg ccc ggg ctg ctc ttt 242
 Gly Ala Gly Ala Gly Glu Ala Ala Leu Pro Leu Pro Gly Leu Leu Phe
 65 70 75

ggc gcc ccc gcg ctg ctg ggc ctg gca ctg gtc ctg gcg ctg gtc ctg 290
 Gly Ala Pro Ala Leu Leu Gly Leu Ala Leu Val Leu Ala Leu Val Leu
 80 85 90 95

gtg ggt ctg gtg agc tgg agg cgg cga cag ccg cgg ctt cgc ggc gcg 338
 Val Gly Leu Val Ser Trp Arg Arg Arg Gln Arg Arg Leu Arg Gly Ala
 100 105 110

tcc tcc gca gag gcc ccc gac gga gac aag gac gcc cca gag ccc ctg 386
 Ser Ser Ala Glu Ala Pro Asp Gly Asp Lys Asp Ala Pro Glu Pro Leu
 115 120 125

gac aag gtc atc att ctg tct ccg gga atc tct gat gcc aca gct cct 434
 Asp Lys Val Ile Ile Leu Ser Pro Gly Ile Ser Asp Ala Thr Ala Pro
 130 135 140

gcc tgg cct cct cct ggg gaa gac cca gga acc acc cca cct ggc cac 482
 Ala Trp Pro Pro Pro Gly Glu Asp Pro Gly Thr Thr Pro Pro Gly His
 145 150 155

agt gtc cct gtg cca gcc aca gag ctg ggc tcc act gaa ctg gtg acc 530
 Ser Val Pro Val Pro Ala Thr Glu Leu Gly Ser Thr Glu Leu Val Thr
 160 165 170 175

acc aag acg gcc ggc cct gag caa caa tag cagggagccg gcaggaggtg 580

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Thr Lys Thr Ala Gly Pro Glu Gln Gln
180

gcccctgccc tccctctgga cccccagcca ggggcttgga aatcaaattc agctcttcac 640
tccagcatgc acatgccttc tttctgggac caggctaacc ctgcagaagc acagacacta 700
cagaccacag cattcagccc ccatggagtt tgggtgtgctt gcctttggct tcagacctca 760
ccatctttga cagcccttga aggtggtagc ccagctcctg ttcctgtgcc ttcaaaaggc 820
tggggcacta tgagtaaaag accgctttta aaatggggaa ggcaccatta agccaaaatg 880
aatctgaaaa aagacaaaa 899

<210> 8
<211> 184
<212> PRT
<213> Homo sapiens

<400> 8

Met Arg Arg Gly Pro Arg Ser Leu Arg Gly Arg Asp Ala Pro Ala Pro
1 5 10 15

Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg His Cys
20 25 30

Val Ala Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro Ala Gly Ala
35 40 45

Ser Ser Pro Ala Pro Arg Thr Ala Leu Gln Pro Gln Glu Ser Val Gly
50 55 60

Ala Gly Ala Gly Glu Ala Ala Leu Pro Leu Pro Gly Leu Leu Phe Gly
65 70 75 80

Ala Pro Ala Leu Leu Gly Leu Ala Leu Val Leu Ala Leu Val Leu Val
85 90 95

Gly Leu Val Ser Trp Arg Arg Arg Gln Arg Arg Leu Arg Gly Ala Ser
100 105 110

Ser Ala Glu Ala Pro Asp Gly Asp Lys Asp Ala Pro Glu Pro Leu Asp
115 120 125

Lys Val Ile Ile Leu Ser Pro Gly Ile Ser Asp Ala Thr Ala Pro Ala
130 135 140

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Trp Pro Pro Pro Gly Glu Asp Pro Gly Thr Thr Pro Pro Gly His Ser
 145 150 155 160

Val Pro Val Pro Ala Thr Glu Leu Gly Ser Thr Glu Leu Val Thr Thr
 165 170 175

Lys Thr Ala Gly Pro Glu Gln Gln
 180

<210> 9
 <211> 26
 <212> PRT
 <213> Homo sapiens

<400> 9

Met Ser Pro Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Leu
 1 5 10 15

Arg His Cys Ile Ala Cys Gly Leu Leu Arg
 20 25

<210> 10
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 10

Val His Val Phe Gly Asp Glu Leu
 1 5

<210> 11
 <211> 34
 <212> PRT
 <213> Homo sapiens

<400> 11

Cys Ser Gln Asn Glu Tyr Phe Asp Ser Leu Leu His Ala Cys Ile Pro
 1 5 10 15

Cys Gln Leu Arg Cys Ser Ser Asn Thr Pro Pro Leu Thr Cys Gln Arg
 20 25 30

Tyr Cys

<210> 12
 <211> 35

12/13

<212> PRT
 <213> Homo sapiens

<400> 12

Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg His Cys Val Ala
 1 5 10 15

Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro Ala Gly Ala Ser Ser
 20 25 30

Pro Ala Pro
 35

<210> 13
 <211> 33
 <212> PRT
 <213> Homo sapiens

<400> 13

Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly Thr Cys Met Ser
 1 5 10 15

Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg Thr Cys Ala Ala Phe
 20 25 30

Cys

<210> 14
 <211> 34
 <212> PRT
 <213> Homo sapiens

<400> 14

Cys Arg Lys Glu Gln Gly Lys Phe Tyr Asp His Leu Leu Arg Asp Cys
 1 5 10 15

Ile Ser Cys Ala Ser Ile Cys Gly Gln His Pro Lys Gln Cys Ala Tyr
 20 25 30

Phe Cys

<210> 15
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 15

Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys Met Asp
1 5 10 15

Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu Gly Cys
20 25 30

<210> 16

<211> 27

<212> PRT

<213> Homo sapiens

<400> 16

Cys His Met Gly Phe Phe Leu Lys Gly Ala Lys Cys Ile Ser Cys His
1 5 10 15

Asp Cys Lys Asn Lys Glu Cys Glu Lys Leu Cys
20 25